# APR 0 7 2004 \$ S

### Sequence Listing

SMITH, VICTORIA

<120> METHODS AND COMPOSITIONS FOR DETECTING DYSPLASIA

<130> P2000R1

<140> US 10/712,124

<141> 2003-11-13

<150> US 60/425,813

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<212> DNA

<213> Homo sapien

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Val Gly Glu Asn Gly Gly Glu Lys Pro Thr Pro Ser Pro Pro Trp 35 40 45

Arg Leu Arg Arg Ser Lys Arg Cys Ser Cys Ser Ser Leu Met Asp
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Lys Glu Cys Val Tyr Phe Cys His Leu Asp Ile Ile Trp Val Asn 65 70 75

Thr Pro Glu His Val Val Pro Tyr Gly Leu Gly Ser Pro Arg Ser 80 85 90

Lys Arg Ala Leu Glu Asn Leu Leu Pro Thr Lys Ala Thr Asp Arg 95 100 105

Glu Asn Arg Cys Gln Cys Ala Ser Gln Lys Asp Lys Lys Cys Trp 110 115 120

Asn Phe Cys Gln Ala Gly Lys Glu Leu Arg Ala Glu Asp Ile Met 125 130 135

Glu Lys Asp Trp Asn Asn His Lys Lys Gly Lys Asp Cys Ser Lys 140 145 150

Leu Gly Lys Lys Cys Ile Tyr Gln Gln Leu Val Arg Gly Arg Lys 155 160 165

Ile Arg Arg Ser Ser Glu Glu His Leu Arg Gln Thr Arg Ser Glu 170 175 180

Thr Met Arg Asn Ser Val Lys Ser Ser Phe His Asp Pro Lys Leu 185 190 195

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His Trp

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Phe Ala Glu Asn Lys Glu Ile Gln Lys Leu Ala Glu Gln Phe Val 95 100 105 Leu Leu Asn Leu Val Tyr Glu Thr Thr Asp Lys His Leu Ser Pro 110 115 Asp Gly Gln Tyr Val Pro Arg Ile Met Phe Val Asp Pro Ser Leu 125 130 135 Thr Val Arg Ala Asp Ile Thr Gly Arg Tyr Ser Asn Arg Leu Tyr

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<213> Homo sapien

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Val Val Leu Pro Arg Arg Leu Pro Gly Pro Arg Val Arg Arg Ala 35 40 45

Leu Pro Ser His Leu Gly Leu His Pro Glu Arg Val Ser Tyr Val
50 55 60

Leu Gly Ala Thr Gly His Asn Phe Thr Leu His Leu Arg Lys Asn 65 70 75

Arg Asp Leu Leu Gly Ser Gly Tyr Thr Glu Thr Tyr Thr Ala Ala 80 85 90

Asn Gly Ser Glu Val Thr Glu Gln Pro Arg Gly Gln Asp His Cys 95 100 105

Leu Tyr Gln Gly His Val Glu Gly Tyr Pro Asp Ser Ala Ala Ser 110 115 120

Leu Ser Thr Cys Ala Gly Leu Arg Gly Phe Phe Gln Val Gly Ser 125 130 135

Asp Leu His Leu Ile Glu Pro Leu Asp Glu Gly Gly Glu Gly Gly 140 145 150

Arg His Ala Val Tyr Gln Ala Glu His Leu Leu Gln Thr Ala Gly
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Thr Cys Gly Val Ser Asp Asp Ser Leu Gly Ser Leu Leu Gly Pro 170 175 180

Arg Thr Ala Ala Val Phe Arg Pro Arg Pro Gly Asp Ser Leu Pro 185 190 195

Ser Arg Glu Thr Arg Tyr Val Glu Leu Tyr Val Val Asp Asn

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Asp	Arg	Phe	His	Val 260	Ser	Pro	Asp	Pro	Ser 265	Val	Thr	Leu	Glu	Asn 270
Leu	Leu	Thr	Trp	Gln 275	Ala	Arg	Gln	Arg	Thr 280	Arg	Arg	His	Leu	His 285
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Arg	Cys	Ile	Met	Ala 365	Gly	Ser	Ile	Gly	Ser 370	Ser	Phe	Pro	Arg	Met 375
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Gln	Cys	Asp	Cys	Gly 425	Pro	Pro	Glu	Asp	Cys 430	Arg	Asn	Arg	Cys	Cys 435
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Cys	Arg	Pro	Lys	Lys 470	Asp	Met	Cys	Asp	Leu 475	Glu	Glu	Phe	Cys	Asp 480
Gly	Arg	His	Pro	Glu 485	Cys	Pro	Glu	Asp	Ala 490	Phe	Gln	Glu	Asn	Gly 495
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Ala Ser	Arg '	Tyr	Arg 545	Ala	Asp	Met	Cys	Gly 550	Val	Leu	Gln	Cys	Lys 555
Gly Gly	Gln	Gln	Pro 560	Leu	Gly	Arg	Ala	Ile 565	Cys	Ile	Val	Asp	Val 570
Cys His	Ala :	Leu	Thr 575	Thr	Glu	Asp	Gly	Thr 580	Ala	Tyr	Glu	Pro	Val 585
Pro Glu	Gly '	Thr	Arg 590	Cys	Gly	Pro	Glu	Lys 595	Val	Cys	Trp	Lys	Gly 600
Arg Cys	Gln A	Asp	Leu 605	His	Val	Tyr	Arg	Ser 610	Ser	Asn	Cys	Ser	Ala 615
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Val Leu	Val 1	Leu	Leu 665	Ala	Val	Val	Leu	Val 670	Thr	Leu	Ala	Gly	Ile 675
Val Leu			665					670				_	675
	Tyr A	Arg Thr	665 Lys 680	Ala	Arg	Ser	Arg	670 Ile 685	Leu	Ser	Arg	Asn	675 Val 690
Ile Val	Tyr i	Arg Thr	665 Lys 680 Thr 695	Ala Met	Arg Gly	Ser Arg	Arg Ser	670 Ile 685 Asn 700	Leu Pro	Ser Leu	Arg Phe	Asn His	675 Val 690 Gln 705
Ile Val	Tyr i	Arg Ihr Arg	665 Lys 680 Thr 695 Val 710	Ala Met Pro	Arg Gly Ala	Ser Arg Lys	Arg Ser Gly	670 Ile 685 Asn 700 Gly 715	Leu Pro Ala	Ser Leu Pro	Arg Phe Ala	Asn His	075 Val 690 Gln 705 Ser 720
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Ile Val Ala Pro Ala Ala Arg Gly	Tyr A	Arg Thr Arg Gln Pro	665 Lys 680 Thr 695 Val 710 Glu 725 Ala 740	Ala Met Pro Leu Ser	Arg Gly Ala Val Ser	Ser Arg Lys Pro	Arg Ser Gly Thr	670 Ile 685 Asn 700 Gly 715 Thr 730 Leu 745	Leu Pro Ala His	Ser Leu Pro Pro	Arg Phe Ala Gly Pro	Asn His Pro Gln	Val 690 Gln 705 Ser 720 Pro 735
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Ile Val Ala Pro Ala Ala Arg Gly Ala Arg Ala Pro	Tyr A Lys : Ser A Pro (	Arg Thr Arg Gln Pro Val Gln Pro	665 Lys 680 Thr 695 Val 710 Glu 725 Ala 740 Thr 755 Ala 770	Ala Met Pro Leu Ser Val	Arg Gly Ala Val Ser Ser	Ser Arg Lys Pro Val Ser	Arg Ser Gly Thr Ala Pro	670 Ile 685 Asn 700 Gly 715 Thr 730 Leu 745 Pro 760 Ile 775	Leu Pro Ala His Lys Phe	Ser Leu Pro Pro Arg Pro	Arg Phe Ala Gly Pro Val	Asn His Pro Gln Pro Pro	Val 690 Gln 705 Ser 720 Pro 735 Pro 750 Val 765

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Thr Gly Gly Ser Ser Ala Val Ala Gly Gln Trp Pro Trp Gln Val 50 55 60

Ser Ile Thr Tyr Glu Gly Val His Val Cys Gly Gly Ser Leu Val
65 70 75

Ser Glu Gln Trp Val Leu Ser Ala Ala His Cys Phe Pro Ser Glu 80 85 90

His His Lys Glu Ala Tyr Glu Val Lys Leu Gly Ala His Gln Leu 95 100 105

Asp Ser Tyr Ser Glu Asp Ala Lys Val Ser Thr Leu Lys Asp Ile 110 115 120

Ile Pro His Pro Ser Tyr Leu Gln Glu Gly Ser Gln Gly Asp Ile 125 130 135

Ala Leu Leu Gln Leu Ser Arg Pro Ile Thr Phe Ser Arg Tyr Ile 140 145 150

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<sup>&</sup>lt;211> 1040

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapien

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Val	Ser	Arg	Glu	Ala 125	Ile	Leu	Arg	Phe	Gly 130	Phe	Leu	Gln	Glu	Phe 135
Ser	Lys	Glu	Glu	Arg 140	Asp	Pro	Val	Lys	Ala 145	His	Glu	Gly	Trp	Gly 150
Val	Met	Leu	Pro	Cys 155	Asn	Pro	Pro	Ala	His 160	Tyr	Pro	Gly	Leu	Ser 165
Tyr	Arg	Trp	Leu	Leu 170	Asn	Glu	Phe	Pro	Asn 175	Phe	Ile	Pro	Thr	Asp 180
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Lys Trp	Asp	Pro	Val 935	Val	Pro	Phe	Arg	Asn 940	Glu	Ser	Ala	Val	Thr 945
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Thr Ser Met Met Val Glu Asn Met Ala Val Arg Pro Ala Pro His 1010 1015 1020

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- Ser Arg Pro Ala Ile Leu Tyr Ala Leu Leu Ser Ser Ser Leu Lys 20 25 30
- Ala Val Pro Arg Pro Arg Ser Arg Cys Leu Cys Arg Gln His Arg 35 40 45
- Pro Val Gln Leu Cys Ala Pro His Arg Thr Cys Arg Glu Ala Leu
  50 55 60
- Asp Val Leu Ala Lys Thr Val Ala Phe Leu Arg Asn Leu Pro Ser
  65 70 75
- Phe Trp Gln Leu Pro Pro Gln Asp Gln Arg Arg Leu Leu Gln Gly 80 85 90
- Cys Trp Gly Pro Leu Phe Leu Leu Gly Leu Ala Gln Asp Ala Val 95 100 105
- Thr Phe Glu Val Ala Glu Ala Pro Val Pro Ser Ile Leu Lys Lys 110 115 120
- Ile Leu Leu Glu Glu Pro Ser Ser Ser Gly Gly Ser Gly Gln Leu 125 130 135
- Pro Asp Arg Pro Gln Pro Ser Leu Ala Ala Val Gln Trp Leu Gln
  140 145 150
- Cys Cys Leu Glu Ser Phe Trp Ser Leu Glu Leu Ser Pro Lys Glu
  155 160 165
- Tyr Ala Cys Leu Lys Gly Thr Ile Leu Phe Asn Pro Asp Val Pro 170 175 180
- Gly Leu Gln Ala Ala Ser His Ile Gly His Leu Gln Gln Glu Ala 185 190 195
- His Trp Val Leu Cys Glu Val Leu Glu Pro Trp Cys Pro Ala Ala 200 205 210
- Gln Gly Arg Leu Thr Arg Val Leu Leu Thr Ala Ser Thr Leu Lys

215 220 225

Ser Ile Pro Thr Ser Leu Leu Gly Asp Leu Phe Phe Arg Pro Ile 230 235 240

Ile Gly Asp Val Asp Ile Ala Gly Leu Leu Gly Asp Met Leu Leu 245 250 250

Leu Arg

<210> 13

<211> 1998

<212> DNA

<213> Homo sapien

<400> 13

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<210> 14

<211> 399

<212> PRT

<213> Homo sapien

### <400> 14

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Met Glu Thr Pro Pro Trp Asp Pro Ala Arg Asn Asp Ser Leu Pro
20 25 30

Pro Thr Leu Thr Pro Ala Val Pro Pro Tyr Val Lys Leu Gly Leu 35 40 45

Thr Val Val Tyr Thr Val Phe Tyr Ala Leu Leu Phe Val Phe Ile 50 55 60

Tyr Val Gln Leu Trp Leu Val Leu Arg Tyr Arg His Lys Arg Leu 65 70 75

Ser Tyr Gln Ser Val Phe Leu Phe Leu Cys Leu Phe Trp Ala Ser 80 85 90

Leu Arg Thr Val Leu Phe Ser Phe Tyr Phe Lys Asp Phe Val Ala

				95					100					105
Ala	Asn	Ser	Leu	Ser 110	Pro	Phe	Val	Phe	Trp 115	Leu	Leu	Tyr	Cys	Phe 120
Pro	Val	Cys	Leu	Gln 125	Phe	Phe	Thr	Leu	Thr 130	Leu	Met	Asn	Leu	Tyr 135
Phe	Thr	Gln	Val	Ile 140	Phe	Lys	Ala	Lys	Ser 145	Lys	Tyr	Ser	Pro	Glu 150
Leu	Leu	Lys	Tyr	Arg 155	Leu	Pro	Leu	Tyr	Leu 160	Ala	Ser	Leu	Phe	Ile 165
Ser	Leu	Val	Phe	Leu 170	Leu	Val	Asn	Leu	Thr 175	Cys	Ala	Val	Leu	Val 180
Lys	Thr	Gly	Asn	Trp 185	Glu	Arg	Lys	Val	Ile 190	Val	Ser	Val	Arg	Val 195
Ala	Ile	Asn	Asp	Thr 200	Leu	Phe	Val	Leu	Cys 205	Ala	Val	Ser	Leu	Ser 210
Ile	Cys	Leu	Tyr	Lys 215	Ile	Ser	Lys	Met	Ser 220	Leu	Ala	Asn	Ile	Tyr 225
Leu	Glu	Ser	Lys	Gly 230	Ser	Ser	Val	Cys	Gln 235	Val	Thr	Ala	Ile	Gly 240
Val	Thr	Val	Ile	Leu 245	Leu	Tyr	Thr	Ser	Arg 250	Ala	Cys	Tyr	Asn	Leu 255
Phe	Ile	Leu	Ser	Phe 260	Ser	Gln	Asn	Lys	Ser 265	Val	His	Ser	Phe	Asp 270
Tyr	Asp	Trp	Tyr	Asn 275	Val	Ser	Asp	Gln	Ala 280	Asp	Leu	Lys	Asn	Gln 285
Leu	Gly	Asp	Ala	Gly 290	Tyr	Val	Leu	Phe	Gly 295	Val	Val	Leu	Phe	Val 300
Trp	Glu	Leu	Leu	Pro 305	Thr	Thr	Leu	Val	Val 310	Tyr	Phe	Phe	Arg	Val 315
Arg	Asn	Pro	Thr	Lys 320	Asp	Leu	Thr	Asn	Pro 325	Gly	Met	Val	Pro	Ser 330
His	Gly	Phe	Ser	Pro 335	Arg	Ser	Tyr	Phe	Phe 340	Asp	Asn	Pro	Arg	Arg 345
Туг	Asp	Ser	Asp	Asp 350	Asp	Leu	Ala	Trp	Asn 355	Ile	Ala	Pro	Gln	Gly 360
Leu	Gln	Gly	Gly	Phe 365	Ala	Pro	Asp	Tyr	Tyr 370	Asp	Trp	Gly	Gln	Gln 375
Thr	Asn	Ser	Phe	Leu 380	Ala	Gln	Ala	Gly	Thr 385	Leu	Gln	Asp	Ser	Thr 390
Leu	Asp	Pro	Asp	Lys 395	Pro	Ser	Leu	Gly						

<210> 15 <211> 2320 <212> DNA <213> Homo sapien

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<210> 16

<211> 509

<212> PRT

<213> Homo sapien

#### <400> 16

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His Phe Asn Arg Ile Ser His Gly Leu Gln Gly Leu Ser Ala Val 20 25 30

Pro Leu Arg Thr Tyr Ala Asp Gln Pro Ile Asp Ala Asp Val Thr
35 40 45

Val Ile Gly Ser Gly Pro Gly Gly Tyr Val Ala Ala Ile Lys Ala 50 55 60

Ala Gln Leu Gly Phe Lys Thr Val Cys Ile Glu Lys Asn Glu Thr 65 70 75

Leu Gly Gly Thr Cys Leu Asn Val Gly Cys Ile Pro Ser Lys Ala Leu Leu Asn Asn Ser His Tyr Tyr His Met Ala His Gly Thr Asp 95 100 105 Phe Ala Ser Arg Gly Ile Glu Met Ser Glu Val Arg Leu Asn Leu Asp Lys Met Met Glu Gln Lys Ser Thr Ala Val Lys Ala Leu Thr 130 Gly Gly Ile Ala His Leu Phe Lys Gln Asn Lys Val Val His Val Asn Gly Tyr Gly Lys Ile Thr Gly Lys Asn Gln Val Thr Ala Thr Lys Ala Asp Gly Gly Thr Gln Val Ile Asp Thr Lys Asn Ile Leu Ile Ala Thr Gly Ser Glu Val Thr Pro Phe Pro Gly Ile Thr Ile Asp Glu Asp Thr Ile Val Ser Ser Thr Gly Ala Leu Ser Leu Lys Lys Val Pro Glu Lys Met Val Val Ile Gly Ala Gly Val Ile Gly Val Glu Leu Gly Ser Val Trp Gln Arg Leu Gly Ala Asp Val Thr Ala Val Glu Phe Leu Gly His Val Gly Gly Val Gly Ile Asp Met Glu Ile Ser Lys Asn Phe Gln Arg Ile Leu Gln Lys Gln Gly Phe 265 Lys Phe Lys Leu Asn Thr Lys Val Thr Gly Ala Thr Lys Lys Ser Asp Gly Lys Ile Asp Val Ser Ile Glu Ala Ala Ser Gly Gly Lys 295 300 Ala Glu Val Ile Thr Cys Asp Val Leu Leu Val Cys Ile Gly Arg 310 Arg Pro Phe Thr Lys Asn Leu Gly Leu Glu Glu Leu Gly Ile Glu 320 325 330 Leu Asp Pro Arg Gly Arg Ile Pro Val Asn Thr Arg Phe Gln Thr Lys Ile Pro Asn Ile Tyr Ala Ile Gly Asp Val Val Ala Gly Pro Met Leu Ala His Lys Ala Glu Asp Glu Gly Ile Ile Cys Val Glu Gly Met Ala Gly Gly Ala Val His Ile Asp Tyr Asn Cys Val Pro

380 385 390 Ser Val Ile Tyr Thr His Pro Glu Val Ala Trp Val Gly Lys Ser 395 400 405 Glu Glu Gln Leu Lys Glu Glu Gly Ile Glu Tyr Lys Val Gly Lys 415 410 Phe Pro Phe Ala Ala Asn Ser Arg Ala Lys Thr Asn Ala Asp Thr Asp Gly Met Val Lys Ile Leu Gly Gln Lys Ser Thr Asp Arg Val 445 450 Leu Gly Ala His Ile Leu Gly Pro Gly Ala Gly Glu Met Val Asn Glu Ala Ala Leu Ala Leu Glu Tyr Gly Ala Ser Cys Glu Asp Ile 475 480 Ala Arg Val Cys His Ala His Pro Thr Leu Ser Glu Ala Phe Arg 490 495 Glu Ala Asn Leu Ala Ala Ser Phe Gly Lys Ser Ile Asn Phe

505

<210> 17

<211> 2090

<212> DNA

<400> 17

<213> Homo sapien

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gggagctgtc ggctggtga ggaggaagtt aacatcccta ataggagggt 200
tctggttact ggtgccactg ggcttcttgg cagagctgta cacaaagaat 250
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ccaaaatttg aacaggttaa tctgttggat tctaatgcag ttcatcacat 350
cattcatgat tttcagcccc atgttatagt acattgtgca gcagaggaga 400
gaccagatgt tgtagaaaat cagccagatg ctgcctctca acttaatgtg 450
gatgcttctg ggaatttagc aaaggaagca gctgctgttg gagcatttct 500
catctacatt agctcagatt atgtatttga tggaacaaat ccaccttaca 550
gagaggaaga cataccagct cccctaaatt tgtatggcaa aacaaaatta 600
gatggagaaa aggctgtcct ggagaacaat ctaggagctg ctgttttgag 650
gattcctatt ctgtatggg aagttgaaaa gctcgaagaa agtgctgtga 700

ctgttatgtt tgataaagtg cagttcagca acaagtcagc aaacatggat 750

cactggcage agaggtteec cacacatgte aaagatgtgg ceactgtgtg 800 ccggcagcta gcagagaaga gaatgctgga tccatcaatt aagggaacct 850 ttcactggtc tggcaatgaa cagatgacta agtatgaaat ggcatgtgca 900 attgcagatg ccttcaacct ccccagcagt cacttaagac ctattactga 950 cagccctgtc ctaggagcac aacgtccgag aaatgctcag cttgactgct 1000 ccaaattgga gaccttgggc attggccaac gaacaccatt tcgaattgga 1050 atcaaagaat cactttggcc tttcctcatt gacaagagat ggagacaaac 1100 ggtctttcat tagtttattt gtgttgggtt ctttttttt tttaaatgaa 1150 aagtatagta tgtggcactt tttaaagaac aaaggaaata gttttgtatg 1200 agtactttaa ttgtgactct taggatcttt caggtaaatg atgctcttgc 1250 actagtgaaa ttgtctaaag aaactaaagg gcagtcatgc cctgtttgca 1300 gtaatttttc tttttatcat tttgtttgtc ctggctaaac ttggagtttg 1350 agtatagtaa attatgatcc ttaaatattt gagagtcagg atgaagcaga 1400 tctgctgtag acttttcaga tgaaattgtt cattctcgta acctccatat 1450 tttcaggatt tttgaagctg ttgacctttt catgttgatt attttaaatt 1500 gtgtgaaata gtataaaaat cattggtgtt cattatttgc tttgcctgag 1550 ctcagatcaa aatgtttgaa gaaaggaact ttatttttgc aagttacgta 1600 cagtttttat gcttgagata tttcaacatg ttatgtatat tggaacttct 1650 acagettgat geeteetget tttatageag tttatgggga geaettgaaa 1700 gagcgtgtgt acatgtattt tttttctagg caaacattga atgcaaacgt 1750 gtatttttt aatataaata tataactgtc cttttcatcc catgttgccg 1800 ctaagtgata tttcatatgt gtggttatac tcataataat gggccttgta 1850 agtettttea ceatteatga ataataataa atatgtaetg etggeatgta 1900 atgcttagtt ttcttgtatt tacttctttt tttaaatgta aggaccaaac 1950 ttctaaacta attgttcttt tgttgcttta atttttaaaa attacattct 2000 tctgatgtaa catgtgatac atacaaaaga atatagttta atatgtattg 2050 

<sup>&</sup>lt;210> 18

<sup>&</sup>lt;211> 334

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 18

Met 1	Val	Gly	Arg	Glu 5	Lys	Glu	Leu	Ser	Ile 10	His	Phe	Val	Pro	Gly 15
Ser	Cys	Arg	Leu	Val 20	Glu	Glu	Glu	Val	Asn 25	Ile	Pro	Asn	Arg	Arg 30
Val	Leu	Val	Thr	Gly 35	Ala	Thr	Gly	Leu	Leu 40	Gly	Arg	Ala	Val	His 45
Lys	Glu	Phe	Gln	Gln 50	Asn	Asn	Trp	His	Ala 55	Val	Gly	Cys	Gly	Phe 60
Arg	Arg	Ala	Arg	Pro 65	Lys	Phe	Glu	Gln	Val 70	Asn	Leu	Leu	Asp	Ser 75
Asn	Ala	Val	His	His 80	Ile	Ile	His	Asp	Phe 85	Gln	Pro	His	Val	Ile 90
Val	His	Cys	Ala	Ala 95	Glu	Arg	Arg	Pro	Asp 100	Val	Val	Glu	Asn	Gln 105
Pro	Asp	Ala	Ala	Ser 110	Gln	Leu	Asn	Val	Asp 115	Ala	Ser	Gly	Asn	Leu 120
Ala	Lys	Glu	Ala	Ala 125	Ala	Val	Gly	Ala	Phe 130	Leu	Ile	Tyr	Ile	Ser 135
Ser	Asp	Tyr	Val	Phe 140	Asp	Gly	Thr	Asn	Pro 145	Pro	Tyr	Arg	Glu	Glu 150
Asp	Ile	Pro	Ala	Pro 155	Leu	Asn	Leu	Tyr	Gly 160	Lys	Thr	Lys	Leu	Asp 165
Gly	Glu	Lys	Ala	Val 170	Leu	Glu	Asn	Asn	Leu 175	Gly	Ala	Ala	Val	Leu 180
Arg	Ile	Pro	Ile	Leu 185	Tyr	Gly	Glu	Val	Glu 190	Lys	Leu	Glu	Glu	Ser 195
Ala	Val	Thr	Val	Met 200	Phe	Asp	Lys	Val	Gln 205	Phe	Ser	Asn	Lys	Ser 210
Ala	Asn	Met	Asp	His 215	Trp	Gln	Gln	Arg	Phe 220	Pro	Thr	His	Val	Lys 225
Asp	Val	Ala	Thr	Val 230	Cys	Arg	Gln	Leu	Ala 235	Glu	Lys	Arg	Met	Leu 240
Asp	Pro	Ser	Ile	Lys 245	Gly	Thr	Phe	His	Trp 250	Ser	Gly	Asn	Glu	Gln 255
Met	Thr	Lys	Tyr	Glu 260	Met	Ala	Cys	Ala	Ile 265	Ala	Asp	Ala	Phe	Asn 270
Leu	Pro	Ser	Ser	His 275	Leu	Arg	Pro	Ile	Thr 280	Asp	Ser	Pro	Val	Leu 285
Gly	Ala	Gln	Arg	Pro 290	Arg	Asn	Ala	Gln	Leu 295	Asp	Cys	Ser	Lys	Leu 300
Glu	Thr	Leu	Gly	Ile	Gly	Gln	Arg	Thr	Pro	Phe	Arg	Ile	Gly	Ile

305 310 315

Lys Glu Ser Leu Trp Pro Phe Leu Ile Asp Lys Arg Trp Arg Gln 320 325 330

Thr Val Phe His

<210> 19

<211> 2380

<212> DNA

<213> Homo sapien

<400> 19

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tgagatggag acccctgggg ccgtggggtc tcaggggtgc ctggtgaatt 1250 ctgcacttac acgtactcaa gggagcgcgc ccgcgttatc ctcgtacctt 1300 tgtcttcttt ccatctgtgg agtcagtggg tgtcggccgc tctgttgtgg 1350 gggaggtgaa ccagggaggg gcagggcaag gcagggcccc cagagctggg 1400 ccacacagtg ggtgctgggc ctcgccccga agcttctggt gcagcagcct 1450 ctggtgctgt ctccgcggaa gtcagggcgg ctggattcca ggacaggagt 1500 gaatgtaaaa ataaatatcg cttagaatgc aggagaaggg tggagaggag 1550 gcaggggccg agggggtgct tggtgccaaa ctgaaattca gtttcttgtg 1600 tggggccttg cggttcagag ctcttggcga gggtggaggg aggagtgtca 1650 tttctatgtg taatttctga gccattgtac tgtctgggct gggggggaca 1700 ctgtccaagg gagtggcccc tatgagttta tattttaacc actgcttcaa 1750 atctcgattt cacttttttt atttatccag ttatatctac atatctgtca 1800 tctaaataaa tggctttcaa acaaagcaac tgggtcatta aaaccagctc 1850 aaagggggtt taaaaaaaaa aaaaccagcc catcetttga ggctgatttt 1900 tctttttttt aagttctatt ttaaaagcta tcaaacagcg acatagccat 1950 acatetgaet geetgaeatg gaeteetgee eacttggggg aaacettata 2000 cccagaggaa aatacacacc tggggagtac atttgacaaa tttcccttag 2050 gatttcgtta tctcaccttg accetcagce aagattggta aagctgcgtc 2100 ctggcgattc caggagaccc agctggaaac ctggcttctc catgtgaggg 2150 gatgggaaag gaaagaagag aatgaagact acttagtaat tcccatcagg 2200 aaatgctgac cttttacata aaatcaagga gactgctgaa aatctctaag 2250 ggacaggatt ttccagatcc taattggaaa tttagcaata aggaggaggag 2300 tccaagggga caaataaagg cagagagaga gagagagaga gggagaggaa 2350 gaaaagagag agagaaaaga gcctcgtgcc 2380

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<210> 20
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Leu Ala Thr Phe Asp Pro Ala Arg Gly Thr Asp Ala Thr Asn Pro  $20 \hspace{1cm} 25 \hspace{1cm} 30$ 

<sup>&</sup>lt;211> 302

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 20

Pro	Glu	Gly	Pro	Gln 35	Asp	Arg	Ser	Ser	Gln 40	Gln	Lys	Gly	Arg	Leu 45
Ser	Leu	Gln	Asn	Thr 50	Ala	Glu	Ile	Gln	His 55	Cys	Leu	Val	Asn	Ala 60
Gly	Asp	Val	Gly	Cys 65	Gly	Val	Phe	Glu	Cys 70	Phe	Glu	Asn	Asn	Ser 75
Cys	Glu	Ile	Arg	Gly 80	Leu	His	Gly	Ile	Cys 85	Met	Thr	Phe	Leu	His 90
Asn	Ala	Gly	Lys	Phe 95	Asp	Ala	Gln	Gly	Lys 100	Ser	Phe	Ile	Lys	Asp 105
Ala	Leu	Lys	Cys	Lys 110	Ala	His	Ala	Leu	Arg 115	His	Arg	Phe	Gly	Cys 120
Ile	Ser	Arg	Lys	Cys 125	Pro	Ala	Ile	Arg	Glu 130	Met	Val	Ser	Gln	Leu 135
Gln	Arg	Glu	Cys	Туг 140	Leu	Lys	His	Asp	Leu 145	Cys	Ala	Ala	Ala	Gln 150
Glu	Asn	Thr	Arg	Val 155	Ile	Val	Glu	Met	Ile 160	His	Phe	Lys	Asp	Leu 165
Leu	Leu	His	Glu	Pro 170	Tyr	Val	Asp	Leu	Val 175	Asn	Leu	Leu	Leu	Thr 180
Cys	Gly	Glu	Glu	Val 185	Lys	Glu	Ala	Ile	Thr 190	His	Ser	Val	Gln	Val 195
Gln	Cys	Glu	Gln	Asn 200	Trp	Gly	Ser	Leu	Cys 205	Ser	Ile	Leu	Ser	Phe 210
Cys	Thr	Ser	Ala	Ile 215	Gln	Lys	Pro	Pro	Thr 220	Ala	Pro	Pro	Glu	Arg 225
Gln	Pro	Gln	Val	Asp 230	Arg	Thr	Lys	Leu	Ser 235	Arg	Ala	His	His	Gly 240
Glu	Ala	Gly	His	His 245	Leu	Pro	Glu	Pro	Ser 250	Ser	Arg	Glu	Thr	Gly 255
Arg	Gly	Ala	Lys	Gly 260	Glu	Arg	Gly	Ser	Lys 265	Ser	His	Pro	Asn	Ala 270
His	Ala	Arg	Gly	Arg	Val	Gly	Gly	Leu	_	Ala	Gln	Gly	Pro	
				275					280					285

Arg Arg

<sup>&</sup>lt;210> 21 <211> 2516 <212> DNA <213> Homo sapien

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<210> 22

<211> 528

<212> PRT

<213> Homo sapien

### <400> 22

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Leu Ser Leu Gly Val Ile Pro Ala Glu Glu Glu Asn Pro Ala Phe
20 25 30

Trp Asn Arg Gln Ala Ala Glu Ala Leu Asp Ala Ala Lys Lys Leu  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Gln Pro Ile Gln Lys Val Ala Lys Asn Leu Ile Leu Phe Leu Gly
50 55 60

Asp	Gly	Leu	Gly	Val 65	Pro	Thr	Val	Thr	Ala 70	Thr	Arg	Ile	Leu	Lys 75
Gly	Gln	Lys	Asn	Gly 80	Lys	Leu	Gly	Pro	Glu 85	Thr	Pro	Leu	Ala	Met 90
Asp	Arg	Phe	Pro	Tyr 95	Leu	Ala	Leu	Ser	Lys 100	Thr	Tyr	Asn	Val	Asp 105
Arg	Gln	Val	Pro	Asp 110	Ser	Ala	Ala	Thr	Ala 115	Thr	Ala	Tyr	Leu	Cys 120
Gly	Val	Lys	Ala	Asn 125	Phe	Gln	Thr	Ile	Gly 130	Leu	Ser	Ala	Ala	Ala 135
Arg	Phe	Asn	Gln	Cys 140	Asn	Thr	Thr	Arg	Gly 145	Asn	Glu	Val	Ile	Ser 150
Val	Met	Asn	Arg	Ala 155	Lys	Gln	Ala	Gly	Lys 160	Ser	Val	Gly	Val	Val 165
Thr	Thr	Thr	Arg	Val 170	Gln	His	Ala	Ser	Pro 175	Ala	Gly	Thr	Tyr	Ala 180
His	Thr	Val	Asn	Arg 185	Asn	Trp	Tyr	Ser	Asp 190	Ala	Asp	Met	Pro	Ala 195
Ser	Ala	Arg	Gln	Glu 200	Gly	Cys	Gln	Asp	Ile 205	Ala	Thr	Gln	Leu	Ile 210
Ser	Asn	Met	Asp	Ile 215	Asp	Val	Ile	Leu	Gly 220	Gly	Gly	Arg	Lys	Tyr 225
Met	Phe	Pro	Met	Gly 230	Thr	Pro	Asp	Pro	Glu 235	Tyr	Pro	Ala	Asp	Ala 240
Ser	Gln	Asn	Gly	Ile 245	Arg	Leu	Asp	Gly	Lys 250	Asn	Leu	Val	Gln	Glu 255
Trp	Leu	Ala	Lys	His 260	Gln	Gly	Ala	Trp	Tyr 265	Val	Trp	Asn	Arg	Thr 270
Glu	Leu	Met	Gln	Ala 275	Ser	Leu	Asp	Gln	Ser 280	Val	Thr	His	Leu	Met 285
Gly	Leu	Phe	Glu	Pro 290	Gly	Asp	Thr	Lys	Tyr 295	Glu	Ile	Leu	Arg	Asp 300
Pro	Thr	Leu	Asp	Pro 305	Ser	Leu	Met	Glu	Met 310	Thr	Glu	Ala	Ala	Leu 315
Arg	Leu	Leu	Ser	Arg 320	Asn	Pro	Arg	Gly	Phe 325	Tyr	Leu	Phe	Val	Glu 330
Gly	Gly	Arg	Ile	Asp 335	His	Gly	His	His	Glu 340	Gly	Val	Ala	Tyr	Gln 345
Ala	Val	Thr	Glu	Ala 350	Val	Met	Phe	Asp	Asp 355	Ala	Ile	Glu	Arg	Ala 360
Gly	Gln	Leu	Thr	Ser	Glu	Glu	Asp	Thr	Leu	Thr	Leu	Val	Thr	Ala

				365					370					375
Asp l	His	Ser	His	Val 380	Phe	Ser	Phe	Gly	Gly 385	Tyr	Thr	Leu	Arg	Gly 390
Ser S	Ser	Ile	Phe	Gly 395	Leu	Ala	Pro	Ser	Lys 400	Ala	Gln	Asp	Ser	Lys 405
Ala	Tyr	Thr	Ser	Ile 410	Leu	Tyr	Gly	Asn	Gly 415	Pro	Gly	Tyr	Val	Phe 420
Asn S	Ser	Gly	Val	Arg 425	Pro	Asp	Val	Asn	Glu 430	Ser	Glu	Ser	Gly	Ser 435
Pro A	Asp	Tyr	Gln	Gln 440	Gln	Ala	Ala	Val	Pro 445	Leu	Ser	Ser	Glu	Thr 450
His (	Gly	Gly	Glu	Asp 455	Val	Ala	Val	Phe	Ala 460	Arg	Gly	Pro	Gln	Ala 465
His I	Leu	Val	His	Gly 470	Val	Gln	Glu	Gln	Ser 475	Phe	Val	Ala	His	Val 480
Met A	Ala	Phe	Ala	Ala 485	Cys	Leu	Glu	Pro	Tyr 490	Thr	Ala	Cys	Asp	Leu 495
Ala I	Leu	Pro	Ala	Cys 500	Thr	Thr	Asp	Ala	Ala 505	His	Pro	Val	Ala	Ala 510
Ser I	Leu	Pro	Leu	Leu 515	Ala	Gly	Thr	Leu	Leu 520	Leu	Leu	Gly	Ala	Ser 525

Ala Ala Pro

<210> 23

<211> 1746

<212> DNA

<213> Homo sapien

<400> 23

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tgacatgctg ctggactgtc gcttccgtgg ccaaccttgt gggcctgaga 550 acttcaccac gatcttcacc cggatgggaa agtgctacac atttaactct 600 ggcgctgatg gggcagagct gctcaccact actaggggtg gcatgggcaa 650 tgggctggac atcatgctgg acgtgcagca ggaggaatat ctacctgtgt 700 ggagggacaa tgaggagacc ccgtttgagg tggggatccg agtgcagatc 750 cacagocagg aggagocgco catcatogat cagotgggot tgggggtgtc 800 cccgggctac cagacetttg tttcttgcca gcagcagcag ctgagettcc 850 tgccaccgcc ctggggcgat tgcagttcag catctctgaa ccccaactat 900 gagccagagc cctctgatcc cctaggctcc cccagcccca gccccagccc 950 tecetatace ettatggggt gtegeetgge etgegaaace egetacgtgg 1000 ctcggaagtg cggctgccga atggtgtaca tgccaggcga cgtgccagtg 1050 tgcagccccc agcagtacaa gaactgtgcc cacccggcca tagatgccat 1100 gettegeaag gaetegtgeg eetgeeecaa eeegtgegee ageaegeget 1150 acgccaagga getetecatg gtgcggatec cgagecgege cgccgcgcgc 1200 ttcctggccc ggaagctcaa ccgcagcgag gcctacatcg cggagaacgt 1250 gctggccctg gacatcttct ttgaggccct caactatgag accgtggagc 1300 agaagaaggc ctatgagatg tcagagctgc ttggtgacat tqggggccag 1350 atggggctgt tcatcggggc cagcctgctc accatcctcg agatcctaga 1400 ctacctctgt gaggtgttcc gagacaaggt cctgggatat ttctggaacc 1450 gacagcactc ccaaaggcac tccagcacca atctgcttca ggaagggctg 1500 ggcagccatc gaacccaagt tccccacctc agcctgggcc ccagacctcc 1550 caccectece tgtgccgtca ccaagactet etcegeetee cacegeacet 1600 gctaccttgt cacacagete tagacetget gtetgtgtee teggageece 1650 gccctgacat cctggacatg cctagcctgc acgtagcttt tccgtcttca 1700 

<sup>&</sup>lt;210> 24

<sup>&</sup>lt;211> 531

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 24

Met Lys Pro Thr Ser Gly Pro Glu Glu Ala Arg Arg Pro Ala Ser  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ 

Asp Ile Arg Val Phe Ala Ser Asn Cys Ser Met His Gly Leu Gly 20 25 30

His	Val	Phe	Gly	Pro 35	Gly	Ser	Leu	Ser	Leu 40	Arg	Arg	Gly	Met	Trp 45
Ala	Ala	Ala	Val	Val 50	Leu	Ser	Val	Ala	Thr 55	Phe	Leu	Tyr	Gln	Val 60
Ala	Glu	Arg	Val	Arg 65	Tyr	Tyr	Arg	Glu	Phe 70	His	His	Gln	Thr	Ala 75
Leu	Asp	Glu	Arg	Glu 80	Ser	His	Arg	Leu	Ile 85	Phe	Pro	Ala	Val	Thr 90
Leu	Cys	Asn	Ile	Asn 95	Pro	Leu	Arg	Arg	Ser 100	Arg	Leu	Thr	Pro	Asn 105
Asp	Leu	His	Trp	Ala 110	Gly	Ser	Ala	Leu	Leu 115	Gly	Leu	Asp	Pro	Ala 120
Glu	His	Ala	Ala	Phe 125	Leu	Arg	Ala	Leu	Gly 130	Arg	Pro	Pro	Ala	Pro 135
Pro	Gly	Phe	Met	Pro 140	Ser	Pro	Thr	Phe	Asp 145	Met	Ala	Gln	Leu	Tyr 150
Ala	Arg	Ala	Gly	His 155	Ser	Leu	Asp	Asp	Met 160	Leu	Leu	Asp	Cys	Arg 165
Phe	Arg	Gly	Gln	Pro 170	Cys	Gly	Pro	Glu	Asn 175	Phe	Thr	Thr	Ile	Phe 180
Thr	Arg	Met	Gly	Lys 185	Cys	Tyr	Thr	Phe	Asn 190	Ser	Gly	Ala	Asp	Gly 195
Ala	Glu	Leu	Leu	Thr 200	Thr	Thr	Arg	Gly	Gly 205	Met	Gly	Asn	Gly	Leu 210
Asp	Ile	Met	Leu	Asp 215	Val	Gln	Gln	Glu	Glu 220	Tyr	Leu	Pro	Val	Trp 225
Arg	Asp	Asn	Glu	Glu 230	Thr	Pro	Phe	Glu	Val 235	Gly	Ile	Arg	Val	Gln 240
Ile	His	Ser	Gln	Glu 245	Glu	Pro	Pro	Ile	Ile 250	Asp	Gln	Leu	Gly	Leu 255
Gly	Val	Ser	Pro	Gly 260	Tyr	Gln	Thr	Phe	Val 265	Ser	Cys	Gln	Gln	Gln 270
Gln	Leu	Ser	Phe	Leu 275	Pro	Pro	Pro	Trp	Gly 280	Asp	Cys	Ser	Ser	Ala 285
Ser	Leu	Asn	Pro	Asn 290	Tyr	Glu	Pro	Glu	Pro 295	Ser	Asp	Pro	Leu	Gly 300
Ser	Pro	Ser	Pro	Ser 305	Pro	Ser	Pro	Pro	Tyr 310	Thr	Leu	Met	Gly	Cys 315
Arg	Leu	Ala	Cys	Glu 320	Thr	Arg	Tyr	Val	Ala 325	Arg	Lys	Cys	Gly	Cys 330

Arg Met Val Tyr Met Pro Gly Asp Val Pro Val Cys Ser Pro Gln 335 Gln Tyr Lys Asn Cys Ala His Pro Ala Ile Asp Ala Met Leu Arq 350 355 360 Lys Asp Ser Cys Ala Cys Pro Asn Pro Cys Ala Ser Thr Arg Tyr Ala Lys Glu Leu Ser Met Val Arg Ile Pro Ser Arg Ala Ala Ala 380 385 390 Arg Phe Leu Ala Arg Lys Leu Asn Arg Ser Glu Ala Tyr Ile Ala 395 400 405 Glu Asn Val Leu Ala Leu Asp Ile Phe Phe Glu Ala Leu Asn Tyr 410 415 420 Glu Thr Val Glu Gln Lys Lys Ala Tyr Glu Met Ser Glu Leu Leu Gly Asp Ile Gly Gln Met Gly Leu Phe Ile Gly Ala Ser Leu Leu Thr Ile Leu Glu Ile Leu Asp Tyr Leu Cys Glu Val Phe Arg 465 Asp Lys Val Leu Gly Tyr Phe Trp Asn Arg Gln His Ser Gln Arg 475 His Ser Ser Thr Asn Leu Leu Gln Glu Gly Leu Gly Ser His Arg 485 490 495 Thr Gln Val Pro His Leu Ser Leu Gly Pro Arg Pro Pro Thr Pro 505 510 Pro Cys Ala Val Thr Lys Thr Leu Ser Ala Ser His Arg Thr Cys 515 520 525 Tyr Leu Val Thr Gln Leu

530

<210> 25

<211> 1104

<212> DNA

<213> Homo sapien

# <400> 25

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gaggaggact gcctgccca taccaggcca aacagttgca cctgcactgg 400 tecgaettge catataaggg eteggageae ageetegatg gggageaett 450 tgccatggag atgcacatag tacatgagaa agagaagggg acatcgagga 500 atgtgaaaga ggcccaggac cctgaagacg aaattgcggt gctggccttt 550 ctggtggagg ctggaaccca ggtgaacgag ggcttccagc cactggtgga 600 ggcactgtct aatatcccca aacctgagat gagcactacg atggcagaga 650 gcagcctgtt ggacctgctc cccaaggagg agaaactgag gcactacttc 700 cgctacctgg gctcactcac cacaccgacc tgcgatgaga aggtcgtctg 750 gactgtgttc cgggagccca ttcagcttca cagagaacag atcctggcat 800 teteteagaa getgtaetae gacaaggaac agacagtgag catgaaggae 850 aatgtcaggc ccctgcagca gctggggcag cgcacggtga taaagtccgg 900 ggcccgggt cggccgctgc cctgggccct qcctgccctg ctgggcccca 950 tgctggcctg cctgctggcc ggcttcctgc gatgatggct cacttctgca 1000 cgcagcctct ctgttgcctc agctctccaa gttccaggct tccggtcctt 1050 agectteeca ggtgggaett taggeatgat taaaatatgg acatattttt 1100 ggag 1104

<210> 26

<211> '312

<212> PRT

<213> Homo sapien

# <400> 26

Met Arg Met Leu Leu Ala Leu Leu Ala Leu Ser Ala Ala Arg Pro 1 5 10 15

Ser Ala Ser Ala Glu Ser His Trp Cys Tyr Glu Val Gln Ala Glu 20 25 30

Ser Ser Asn Tyr Pro Cys Leu Val Pro Val Lys Trp Gly Gly Asn 35 40 45

Cys Gln Lys Asp Arg Gln Ser Pro Ile Asn Ile Val Thr Thr Lys
50 55 60

Ala Lys Val Asp Lys Lys Leu Gly Arg Phe Phe Ser Gly Tyr
65 70 75

Asp Lys Gln Thr Trp Thr Val Gln Asn Asn Gly His Ser Val

Met Met Leu Leu Glu Asn Lys Ala Ser Ile Ser Gly Gly Leu 95 100 105

Pro Ala Pro Tyr Gln Ala Lys Gln Leu His Leu His Trp Ser Asp

110 115 120 Leu Pro Tyr Lys Gly Ser Glu His Ser Leu Asp Gly Glu His Phe 130 Ala Met Glu Met His Ile Val His Glu Lys Glu Lys Gly Thr Ser 140 145 150 Arg Asn Val Lys Glu Ala Gln Asp Pro Glu Asp Glu Ile Ala Val Leu Ala Phe Leu Val Glu Ala Gly Thr Gln Val Asn Glu Gly Phe 175 180 Gln Pro Leu Val Glu Ala Leu Ser Asn Ile Pro Lys Pro Glu Met 185 190 195 Ser Thr Thr Met Ala Glu Ser Ser Leu Leu Asp Leu Leu Pro Lys 205 210 Glu Glu Lys Leu Arg His Tyr Phe Arg Tyr Leu Gly Ser Leu Thr 215 Thr Pro Thr Cys Asp Glu Lys Val Val Trp Thr Val Phe Arg Glu 230 235 240 Pro Ile Gln Leu His Arq Glu Gln Ile Leu Ala Phe Ser Gln Lys 255 Leu Tyr Tyr Asp Lys Glu Gln Thr Val Ser Met Lys Asp Asn Val 265 Arg Pro Leu Gln Gln Leu Gly Gln Arg Thr Val Ile Lys Ser Gly 280 285 Ala Pro Gly Arg Pro Leu Pro Trp Ala Leu Pro Ala Leu Leu Gly 295 300 Pro Met Leu Ala Cys Leu Leu Ala Gly Phe Leu Arg 310

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<211> 585

<212> DNA

<213> Homo sapien

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<211> 148

<212> PRT

<213> Homo sapien

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Asp Ser Gly Ile Ser Pro Arg Ala Val Trp Gln Phe Arg Lys Met 20 25 30

Ile Lys Cys Val Ile Pro Gly Ser Asp Pro Phe Leu Glu Tyr Asn
35 40 45

Asn Tyr Gly Cys Tyr Cys Gly Leu Gly Gly Ser Gly Thr Pro Val
50 55 60

Asp Glu Leu Asp Lys Cys Cys Gln Thr His Asp Asn Cys Tyr Asp 65 70 75

Gln Ala Lys Lys Leu Asp Ser Cys Lys Phe Leu Leu Asp Asn Pro 80 85 90

Tyr Thr His Thr Tyr Ser Tyr Ser Cys Ser Gly Ser Ala Ile Thr 95 100 105

Cys Ser Ser Lys Asn Lys Glu Cys Glu Ala Phe Ile Cys Asn Cys 110 115 120

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Ala His Lys Asn Leu Asp Thr Lys Lys Tyr Cys Gln Ser 140 145

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<211> 397

<212> PRT

<213> Homo sapien

<400> 30

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Leu Ala Ala Ser Leu Ser Cys Ser Gly Thr Ile Gln Gly Thr Asn 20 25 30

Arg Ser Ser Lys Gly Arg Ser Leu Ile Gly Lys Val Asp Gly Thr
35 40 45

Ser	His	Val	Thr	Gly 50	Lys	Gly	Val	Thr	Val 55	Glu	Thr	Val	Phe	Ser 60
Val	Asp	Glu	Phe	Ser 65	Ala	Ser	Val	Leu	Thr 70	Gly	Lys	Leu	Thr	Thr 75
Val	Phe	Leu	Pro	Ile 80	Val	Tyr	Thr	Ile	Val 85	Phe	Val	Val	Gly	Leu 90
Pro	Ser	Asn	Gly	Met 95	Ala	Leu	Trp	Val	Phe 100	Leu	Phe	Arg	Thr	Lys 105
Lys	Lys	His	Pro	Ala 110	Val	Ile	Tyr	Met	Ala 115	Asn	Leu	Ala	Leu	Ala 120
Asp	Leu	Leu	Ser	Val 125	Ile	Trp	Phe	Pro	Leu 130	Lys	Ile	Ala	Tyr	His 135
Ile	His	Gly	Asn	Asn 140	Trp	Ile	Tyr	Gly	Glu 145	Ala	Leu	Cys	Asn	Val 150
Leu	Ile	Gly	Phe	Phe 155	Tyr	Gly	Asn	Met	Tyr 160	Cys	Ser	Ile	Leu	Phe 165
Met	Thr	Cys	Leu	Ser 170	Val	Gln	Arg	Tyr	Trp 175	Val	Ile	Val	Asn	Pro 180
Met	Gly	His	Ser	Arg 185	Lys	Lys	Ala	Asn	Ile 190	Ala	Ile	Gly	Ile	Ser 195
Leu	Ala	Ile	Trp	Leu 200	Leu	Ile	Leu	Leu	Val 205	Thr	Ile	Pro	Leu	Tyr 210
Val	Val	Lys	Gln	Thr 215	Ile	Phe	Ile	Pro	Ala 220	Leu	Asn	Ile	Thr	Thr 225
Cys	His	Asp	Val	Leu 230	Pro	Glu	Gln	Leu	Leu 235	Val	Gly	Asp	Met	Phe 240
Asn	Tyr	Phe	Leu	Ser 245	Leu	Ala	Ile	Gly	Val 250	Phe	Leu	Phe	Pro	Ala 255
Phe	Leu	Thr	Ala	Ser 260	Ala	Tyr	Val	Leu	Met 265	Ile	Arg	Met	Leu	Arg 270
Ser	Ser	Ala	Met	Asp 275	Glu	Asn	Ser	Glu	Lys 280	Lys	Arg	Lys	Arg	Ala 285
Ile	Lys	Leu	Ile	Val 290	Thr	Val	Leu	Ala	Met 295	Tyr	Leu	Ile	Cys	Phe 300
Thr	Pro	Ser	Asn	Leu 305	Leu	Leu	Val	Val	His 310	Tyr	Phe	Leu	Ile	Lys 315
Ser	Gln	Gly	Gln	Ser 320	His	Val	Tyr	Ala	Leu 325	Tyr	Ile	Val	Ala	Leu 330
Cys	Leu	Ser	Thr	Leu 335	Asn	Ser	Cys	Ile	Asp 340	Pro	Phe	Val	Tyr	Tyr 345
Phe	Val	Ser	His	Asp	Phe	Arg	Asp	His	Ala	Lys	Asn	Ala	Leu	Leu

350 355 360

Cys Arg Ser Val Arg Thr Val Lys Gln Met Gln Val Ser Leu Thr 365 370 375

Ser Lys Lys His Ser Arg Lys Ser Ser Ser Tyr Ser Ser Ser Ser Ser 380 385 390

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<212> DNA

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<211> 1019

<212> PRT

<213> Homo sapien

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Leu Cys Gly Phe Gln Lys Lys Thr Tyr Ser Lys Met Asn Asn Pro 35 40 45

Ala Ile Lys Arg Ile Gly Asn His Ile Thr Lys Ser Pro Glu Asp
50 55 60

Lys Arg Glu Tyr Arg Gly Leu Glu Leu Ala Asn Gly Ile Lys Val 65 70 75

Leu Leu Met Ser Asp Pro Thr Thr Asp Lys Ser Ser Ala Ala Leu 80 85 90

Asp Val His Ile Gly Ser Leu Ser Asp Pro Pro Asn Ile Ala Gly 95 100 105

Leu Ser His Phe Cys Glu His Met Leu Phe Leu Gly Thr Lys Lys 110 115 120

Tyr Pro Lys Glu Asn Glu Tyr Ser Gln Phe Leu Ser Glu His Ala 125 130 135

Gly Ser Ser Asn Ala Phe Thr Ser Gly Glu His Thr Asn Tyr Tyr

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Ala	Gln	Phe	Phe	Leu 170	Cys	Pro	Leu	Phe	Asp 175	Glu	Ser	Cys	Lys	Asp 180
Arg	Glu	Val	Asn	Ala 185	Val	Asp	Ser	Glu	His 190	Glu	Lys	Asn	Val	Met 195
Asn	Asp	Ala	Trp	Arg 200	Leu	Phe	Gln	Leu	Glu 205	Lys	Ala	Thr	Gly	Asn 210
Pro	Lys	His	Pro	Phe 215	Ser	Lys	Phe	Gly	Thr 220	Gly	Asn	Lys	Tyr	Thr 225
Leu	Glu	Thr	Arg	Pro 230	Asn	Gln	Glu	Gly	Ile 235	Asp	Val	Arg	Gln	Glu 240
Leu	Leu	Lys	Phe	His 245	Ser	Ala	Tyr	Tyr	Ser 250	Ser	Asn	Leu	Met	Ala 255
Val	Cys	Val	Leu	Gly 260	Arg	Glu	Ser	Leu	Asp 265	Asp	Leu	Thr	Asn	Leu 270
Val	Val	Lys	Leu	Phe 275	Ser	Glu	Val	Glu	Asn 280	Lys	Asn	Val	Pro	Leu 285
Pro	Glu	Phe	Pro	Glu 290	His	Pro	Phe	Gln	Glu 295	Glu	His	Leu	Lys	Gln 300
Leu	Tyr	Lys	Ile	Val 305	Pro	Ile	Lys	Asp	Ile 310	Arg	Asn	Leu	Tyr	Val 315
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Gly	His	Tyr	Leu	Gly 335	His	Leu	Ile	Gly	His 340	Glu	Gly	Pro	Gly	Ser 345
Leu	Leu	Ser	Glu	Leu 350	Lys	Ser	Lys	Gly	Trp 355	Val	Asn	Thr	Leu	Val 360
Gly	Gly	Gln	Lys	Glu 365	Gly	Ala	Arg	Gly	Phe 370	Met	Phe	Phe	Ile	Ile 375
Asn	Val	Asp	Leu	Thr 380	Glu	Glu	Gly	Leu	Leu 385	His	Val	Glu	Asp	Ile 390
Ile	Leu	His	Met	Phe 395	Gln	Tyr	Ile	Gln	Lys 400	Leu	Arg	Ala	Glu	Gly 405
Pro	Gln	Glu	Trp	Val 410	Phe	Gln	Glu	Cys	Lys 415	Asp	Leu	Asn	Ala	Val 420
Ala	Phe	Arg	Phe	Lys 425	Asp	Lys	Glu	Arg	Pro 430	Arg	Gly	Tyr	Thr	Ser 435
Lys	Ile	Ala	Gly	Ile 440	Leu	His	Tyr	Tyr	Pro 445	Leu	Glu	Glu	Val	Leu 450

Thr	Ala	Glu	Tyr	Leu 455	Leu	Glu	Glu	Phe	Arg 460	Pro	Asp	Leu	Ile	Glu 465
Met	Val	Leu	Asp	Lys 470	Leu	Arg	Pro	Glu	Asn 475	Val	Arg	Val	Ala	Ile 480
Val	Ser	Lys	Ser	Phe 485	Glu	Gly	Lys	Thr	Asp 490	Arg	Thr	Glu	Glu	Trp 495
Tyr	Gly	Thr	Gln	Tyr 500	Lys	Gln	Glu	Ala	Ile 505	Pro	Asp	Glu	Val	Ile 510
Lys	Lys	Trp	Gln	Asn 515	Ala	Asp	Leu	Asn	Gly 520	Lys	Phe	Lys	Leu	Pro 525
Thr	Lys	Asn	Glu	Phe 530	Ile	Pro	Thr	Asn	Phe 535	Glu	Ile	Leu	Pro	Leu 540
Glu	Lys	Glu	Ala	Thr 545	Pro	Tyr	Pro	Ala	Leu 550	Ile	Lys	Asp	Thr	Val 555
Met	Ser	Lys	Leu	Trp 560	Phe	Lys	Gln	Asp	Asp 565	Lys	Lys	Lys	Lys	Pro 570
Lys	Ala	Cys	Leu	Asn 575	Phe	Glu	Phe	Phe	Ser 580	Pro	Phe	Ala	Tyr	Val 585
Asp	Pro	Leu	His	Cys 590	Asn	Met	Ala	Tyr	Leu 595	Tyr	Leu	Glu	Leu	Leu 600
Lys	Asp	Ser	Leu	Asn 605	Glu	Tyr	Ala	Tyr	Ala 610	Ala	Glu	Leu	Ala	Gly 615
Leu	Ser	Tyr	Asp	Leu 620	Gln	Asn	Thr	Ile	Tyr 625	Gly	Met	Tyr	Leu	Ser 630
Val	Lys	Gly	Tyr	Asn 635	Asp	Lys	Gln	Pro	Ile 640	Leu	Leu	Lys	Lys	Ile 645
Ile	Glu	Lys	Met	Ala 650	Thr	Phe	Glu	Ile	Asp 655	Glu	Lys	Arg	Phe	Glu 660
Ile	Ile	Lys	Glu	Ala 665	Tyr	Met	Arg	Ser	Leu 670	Asn	Asn	Phe	Arg	Ala 675
Glu	Gln	Pro	His	Gln 680	His	Ala	Met	Tyr	Tyr 685	Leu	Arg	Leu	Leu	Met 690
Thr	Glu	Val	Ala	Trp 695	Thr	Lys	Asp	Glu	Leu 700	Lys	Glu	Ala	Leu	Asp 705
Asp	Val	Thr	Leu	Pro 710	Arg	Leu	Lys	Ala	Phe 715	Ile	Pro	Gln	Leu	Leu 720
Ser	Arg	Leu	His	Ile 725	Glu	Ala	Leu	Leu	His 730	Gly	Asn	Ile	Thr	Lys 735
Gln	Ala	Ala	Leu	Gly 740	Ile	Met	Gln	Met	Val 745	Glu	Asp	Thr	Leu	Ile 750

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Glu His Ala His Thr Lys Pro Leu Leu Pro Ser Gln Leu Val Arq
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Tyr Arg Glu Val Gln Leu Pro Asp Arg Gly Trp Phe Val Tyr Gln
                770
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Gln Arg Asn Glu Val His Asn Asn Cys Gly Ile Glu Ile Tyr Tyr
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Gln Thr Asp Met Gln Ser Thr Ser Glu Asn Met Phe Leu Glu Leu
                                     805
                                                         810
Phe Cys Gln Ile Ile Ser Glu Pro Cys Phe Asn Thr Leu Arg Thr
                                     820
Lys Glu Gln Leu Gly Tyr Ile Val Phe Ser Gly Pro Arg Arg Ala
                                     835
Asn Gly Ile Gln Ser Leu Arg Phe Ile Ile Gln Ser Glu Lys Pro
                                     850
Pro His Tyr Leu Glu Ser Arg Val Glu Ala Phe Leu Ile Thr Met
                                     865
Glu Lys Ser Ile Glu Asp Met Thr Glu Glu Ala Phe Gln Lys His
                                     880
Ile Gln Ala Leu Ala Ile Arg Arg Leu Asp Lys Pro Lys Lys Leu
Ser Ala Glu Cys Ala Lys Tyr Trp Gly Glu Ile Ile Ser Gln Gln
                                                         915
Tyr Asn Phe Asp Arg Asp Asn Thr Glu Val Ala Tyr Leu Lys Thr
Leu Thr Lys Glu Asp Ile Ile Lys Phe Tyr Lys Glu Met Leu Ala
                                                         945
Val Asp Ala Pro Arg Arg His Lys Val Ser Val His Val Leu Ala
                                     955
Arg Glu Met Asp Ser Cys Pro Val Val Gly Glu Phe Pro Cys Gln
                                     970
                                                         975
Asn Asp Ile Asn Leu Ser Gln Ala Pro Ala Leu Pro Gln Pro Glu
                980
                                     985
Val Ile Gln Asn Met Thr Glu Phe Lys Arg Gly Leu Pro Leu Phe
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Pro Leu Val Lys Pro His Ile Asn Phe Met Ala Ala Lys Leu
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<sup>&</sup>lt;211> 3624

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 33

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<210> 34

<211> 1043

<212> PRT

<213> Homo sapien

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Leu Glu Pro Leu Val Glu Glu Ser Leu Leu Lys Asn Leu Gln Leu 20 25 30

Arg Tyr Glu Asn Lys Glu Ile Tyr Thr Tyr Ile Gly Asn Val Val 35 40 45

Ile Ser Val Asn Pro Tyr Gln Gln Leu Pro Ile Tyr Gly Pro Glu
50 55 60

e mile .

Phe Ile Ala Lys Tyr Gln Asp Tyr Thr Phe Tyr Glu Leu Lys Pro 65 70 75

His Ile Tyr Ala Leu Ala Asn Val Ala Tyr Gln Ser Leu Arg Asp 80 85 90

Arg Asp Arg Asp Gln Cys Ile Leu Ile Thr Gly Glu Ser Gly Ser  $95 \hspace{1.5cm} 100 \hspace{1.5cm} 105$ 

Gly Lys Thr Glu Ala Ser Lys Leu Val Met Ser Tyr Val Ala Ala 110 115 120

Val Cys Gly Lys Gly Glu Gln Val Asn Ser Val Lys Glu Gln Leu 125 130 135

Leu Gl<br/>n Ser Asn Pro Val Leu Glu Ala Phe Gly Asn Ala Lys Th<br/>r 140 145 150

Ile Arg Asn Asn Asn Ser Ser Arg Phe Gly Lys Tyr Met Asp Ile 155 160 165

Glu	Phe	Asp	Phe	Lys 170	Gly	Ser	Pro	Leu	Gly 175	Gly	Val	Ile	Thr	Asn 180
Tyr	Leu	Leu	Glu	Lys 185	Ser	Arg	Leu	Val	Lys 190	Gln	Leu	Lys	Gly	Glu 195
Arg	Asn	Phe	His	Ile 200	Phe	Tyr	Gln	Leu	Leu 205	Ala	Gly	Ala	Asp	Glu 210
Gln	Leu	Leu	Lys	Ala 215	Leu	Lys	Leu	Glu	Arg 220	Asp	Thr	Thr	Gly	Tyr 225
Ala	Tyr	Leu	Asn	His 230	Glu	Val	Ser	Arg	Val 235	Asp	Gly	Met	Asp	Asp 240
Ala	Ser	Ser	Phe	Arg 245	Ala	Val	Gln	Ser	Ala 250	Met	Ala	Val	Ile	Gly 255
Phe	Ser	Glu	Glu	Glu 260	Ile	Arg	Gln	Val	Leu 265	Glu	Val	Thr	Ser	Met 270
Val	Leu	Lys	Leu	Gly 275	Asn	Val	Leu	Val	Ala 280	Asp	Glu	Phe	Gln	Ala 285
Ser	Gly	Ile	Pro	Ala 290	Ser	Gly	Ile	Arg	Asp 295	Gly	Arg	Gly	Val	Arg 300
Glu	Ile	Gly	Glu	Met 305	Val	Gly	Leu	Asn	Ser 310	Glu	Glu	Val	Glu	Arg 315
Ala	Leu	Cys	Ser	Arg 320	Thr	Met	Glu	Thr	Ala 325	Lys	Glu	Lys	Val	Val 330
Thr	Ala	Leu	Asn	Val 335	Met	Gln	Ala	Gln	Tyr 340	Ala	Arg	Asp	Ala	Leu 345
Ala	Lys	Asn	Ile	Tyr 350	Ser	Arg	Leu	Phe	Asp 355	Trp	Ile	Val	Asn	Arg 360
Ile	Asn	Glu	Ser	Ile 365	Lys	Val	Gly	Ile	Gly 370	Glu	Lys	Lys	Lys	Val 375
Met	Gly	Val	Leu	Asp 380	Ile	Tyr	Gly	Phe	Glu 385	Ile	Leu	Glu	Asp	Asn 390
Ser	Phe	Glu	Gln	Phe 395	Val	Ile	Asn	Tyr	Cys 400	Asn	Glu	Lys	Leu	Gln 405
Gln	Val	Phe	Ile	Glu 410	Met	Thr	Leu	Lys	Glu 415	Glu	Gln	Glu	Glu	Tyr 420
Lys	Arg	Glu	Gly	Ile 425	Pro	Trp	Thr	Lys	Val 430	Asp	Tyr	Phe	Asp	Asn 435
Gly	Ile	Ile	Cys	Lys 440	Leu	Ile	Glu	His	Asn 445	Gln	Arg	Gly	Ile	Leu 450
Ala	Met	Leu	Asp	Glu 455	Glu	Cys	Leu	Arg	Pro 460	Gly	Val	Val	Ser	Asp 465

Ser	Thr	Phe	Leu	Ala 470	Lys	Leu	Asn	Gln	Leu 475	Phe	Ser	Lys	His	Gly 480
His	Tyr	Glu	Ser	Lys 485	Val	Thr	Gln	Asn	Ala 490	Gln	Arg	Gln	Tyr	Asp 495
His	Thr	Met	Gly	Leu 500	Ser	Cys	Phe	Arg	Ile 505	Cys	His	Tyr	Ala	Gly 510
Lys	Val	Thr	Tyr	Asn 515	Val	Thr	Ser	Phe	Ile 520	Asp	Lys	Asn	Asn	Asp 525
Leu	Leu	Phe	Arg	Asp 530	Leu	Leu	Gln	Ala	Met 535	Trp	Lys	Ala	Gln	His 540
Pro	Leu	Leu	Arg	Ser 545	Leu	Phe	Pro	Glu	Gly 550	Asn	Pro	Lys	Gln	Ala 555
Ser	Leu	Lys	Arg	Pro 560	Pro	Thr	Ala	Gly	Ala 565	Gln	Phe	Lys	Ser	Ser 570
Val	Ala	Ile	Leu	Met 575	Lys	Asn	Leu	Tyr	Ser 580	Lys	Ser	Pro	Asn	Tyr 585
Ile	Arg	Cys	Ile	Lys 590	Pro	Asn	Glu	His	Gln 595	Gln	Arg	Gly	Gln	Phe 600
Ser	Ser	Asp	Leu	Val 605	Ala	Thr	Gln	Ala	Arg 610	Tyr	Leu	Gly	Leu	Leu 615
Glu	Asn	Val	Arg	Val 620	Arg	Arg	Ala	Gly	Tyr 625	Ala	His	Arg	Gln	Gly 630
Tyr	Gly	Pro	Phe	Leu 635	Glu	Arg	Tyr	Arg	Leu 640	Leu	Ser	Arg	Ser	Thr 645
Trp	Pro	His	Trp	Asn 650	Gly	Gly	Asp	Arg	Glu 655	Gly	Val	Glu	Lys	Val 660
Leu	Gly	Glu	Leu	Ser 665	Met	Ser	Ser	Gly	Glu 670	Leu	Ala	Phe	Gly	Lys 675
Thr	Lys	Ile	Phe	Ile 680	Arg	Ser	Pro	Lys	Thr 685	Leu	Phe	Tyr	Leu	Glu 690
Glu	Gln	Arg	Arg	Leu 695	Arg	Leu	Gln	Gln	Leu 700	Ala	Thr	Leu	Ile	Gln 705
Lys	Ile	Tyr	Arg	Gly 710	Trp	Arg	Cys	Arg	Thr 715	His	Tyr	Gln	Leu	Met 720
Arg	Lys	Ser	Gln	Ile 725	Leu	Ile	Ser	Ser	Trp 730	Phe	Arg	Gly	Asn	Met 735
Gln	Lys	Lys	Cys	Tyr 740	Gly	Lys	Ile	Lys	Ala 745	Ser	Val	Leu	Leu	Ile 750
Gln	Ala	Phe	Val	Arg 755	Gly	Trp	Lys	Ala	Arg 760	Lys	Asn	Tyr	Arg	Lys 765
Tyr	Phe	Arg	Ser	Glu	Ala	Ala	Leu	Thr	Leu	Ala	Asp	Phe	Ile	Tyr

				770					775					780
Lys S	er 1	Met	Val	Gln 785	Lys	Phe	Leu	Leu	Gly 790	Leu	Lys	Asn	Asn	Leu 795
Pro Se	er '	Thr	Asn	Val 800	Leù	Asp	Lys	Thr	Trp 805	Pro	Ala	Ala	Pro	Tyr 810
Lys C	ys 1	Leu	Ser	Thr 815	Ala	Asn	Gln	Glu	Leu 820	Gln	Gln	Leu	Phe	Tyr 825
Gln Tı	rp 1	Lys	Cys	Lys 830	Arg	Phe	Arg	Asp	Gln 835	Leu	Ser	Pro	Lys	Gln 840
Val G	lu I	Ile	Leu	Arg 845	Glu	Lys	Leu	Cys	Ala 850	Ser	Glu	Leu	Phe	Lys 855
Gly Ly	ys 1	Lys	Ala	Ser 860	Tyr	Pro	Gln	Ser	Val 865	Pro	Ile	Pro	Phe	Cys 870
Gly As	sp :	ľyr	Ile	Gly 875	Leu	Gln	Gly	Asn	Pro 880	Lys	Leu	Gln	Lys	Leu 885
Lys G	ly (	Gly	Glu	Glu 890	Gly	Pro	Val	Leu	Met 895	Ala	Glu	Ala	Val	Lys 900
Lys Va	al <i>P</i>	Asn	Arg	Gly 905	Asn	Gly	Lys	Thr	Ser 910	Ser	Arg	Ile	Leu	Leu 915
Leu Th	nr I	Lys	Gly	His 920	Val	Ile	Leu	Thr	Asp 925	Thr	Lys	Lys	Ser	Gln 930
Ala Ly	ys ]	Ile	Val	Ile 935	Gly	Leu	Asp	Asn	Val 940	Ala	Gly	Val	Ser	Val 945
Thr Se	er I	Leu	Lys	Asp 950	Gly	Leu	Phe	Ser	Leu 955	His	Leu	Ser	Glu	Met 960
Ser Se	er (	/al	Gly	Ser 965	Lys	Gly	Asp	Phe	Leu 970	Leu	Val	Ser	Glu	His 975
Val Il	le G	Glu	Leu	Leu 980	Thr	Lys	Met	Tyr	Arg 985	Ala	Val	Leu	Asp	Ala 990
Thr Gl	ln <i>F</i>	Arg	Gln	Leu 995	Thr	Val	Thr		Thr .000	Glu	Lys	Phe		Val .005
Arg Ph	ne I	.ys		Asn 010	Ser	Val	Ala		Lys .015	Val	Val	Gln		Pro .020
Ala Gl	Ly G	Sly		Asn 025	Ser	Lys	Leu		Tyr 030	Lys	Lys	Lys		Ser .035
His Cy	/s I	Leu		Val 040	Thr	Val	Gln							

<210> 35 <211> 1876 <212> DNA <213> Homo sapien

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<210> 36

<211> 502

<212> PRT

<213> Homo sapien

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- Ala Ala Asp Phe Leu Lys Arg Arg Pro Lys Asn Tyr Pro Pro 35 40 45
- Gly Pro Trp Arg Leu Pro Phe Leu Gly Asn Phe Phe Leu Val Asp
  50 55 60
- Phe Glu Gln Ser His Leu Glu Val Gln Leu Phe Val Lys Lys Tyr
  65 70 75
- Ile Thr Gly Leu Pro Leu Ile Lys Glu Ala Leu Ile His Met Asp 95 100 105
- Gln Asn Phe Gly Asn Arg Pro Val Thr Pro Met Arg Glu His Ile 110 115 120
- Phe Lys Lys Asn Gly Leu Ile Met Ser Ser Gly Gln Ala Trp Lys 125 130 135
- Glu Gln Arg Arg Phe Thr Leu Thr Ala Leu Arg Asn Phe Gly Leu 140 145 150
- Gly Lys Lys Ser Leu Glu Glu Arg Ile Gln Glu Glu Ala Gln His 155 160 165
- Leu Thr Glu Ala Ile Lys Glu Glu Asn Gly Gln Pro Phe Asp Pro
  170 175 180
- His Phe Lys Ile Asn Asn Ala Val Ser Asn Ile Ile Cys Ser Ile 185 \$190\$

Thr	Phe	Gly	Glu	Arg 200	Phe	Glu	Tyr	Gln	Asp 205	Ser	Trp	Phe	Gln	Gln 210
Leu	Leu	Lys	Leu	Leu 215	Asp	Glu	Val	Thr	Tyr 220	Leu	Glu	Ala	Ser	Lys 225
Thr	Cys	Gln	Leu	Tyr 230	Asn	Val	Phe	Pro	Trp 235	Ile	Met	Lys	Phe	Leu 240
Pro	Gly	Pro	His	Gln 245	Thr	Leu	Phe	Ser	Asn 250	Trp	Lys	Lys	Leu	Lys 255
Leu	Phe	Val	Ser	His 260	Met	Ile	Asp	Lys	His 265	Arg	Lys	Asp	Trp	Asn 270
Pro	Ala	Glu	Thr	Arg 275	Asp	Phe	Ile	Asp	Ala 280	Tyr	Leu	Lys	Glu	Met 285
Ser	Lys	His	Thr	Gly 290	Asn	Pro	Thr	Ser	Ser 295	Phe	His	Glu	Glu	Asn 300
Leu	Ile	Cys	Ser	Thr 305	Leu	Asp	Leu	Phe	Phe 310	Ala	Gly	Thr	Glu	Thr 315
Thr	Ser	Thr	Thr	Leu 320	Arg	Trp	Ala	Leu	Leu 325	Tyr	Met	Ala	Leu	Tyr 330
Pro	Glu	Ile	Gln	Glu 335	Lys	Val	Gln	Ala	Glu 340	Ile	Asp	Arg	Val	Ile 345
Gly	Gln	Gly	Gln	Gln 350	Pro	Ser	Thr	Ala	Ala 355	Arg	Glu	Ser	Met	Pro 360
Tyr	Thr	Asn	Ala	Val 365	Ile	His	Glu	Val	Gln 370	Arg	Met	Gly	Asn	Ile 375
Ile	Pro	Leu	Asn	Val 380	Pro	Arg	Glu	Val	Thr 385	Val	Asp	Thr	Thr	Leu 390
Ala	Gly	Tyr	His	Leu 395	Pro	Lys	Gly	Thr	Met 400	Ile	Leu	Thr	Asn	Leu 405
Thr	Ala	Leu	His	Arg 410	Asp	Pro	Thr	Glu	Trp 415	Ala	Thr	Pro	Asp	Thr 420
Phe	Asn	Pro	Asp	His 425	Phe	Leu	Glu	Asn	Gly 430	Gln	Phe	Lys	Lys	Arg 435
Glu	Ala	Phe	Met	Pro 440	Phe	Ser	Ile	Gly	Lys 445	Arg	Ala	Cys	Leu	Gly 450
Glu	Gln	Leu	Ala	Arg 455	Thr	Glu	Leu	Phe	Ile 460	Phe	Phe	Thr	Ser	Leu 465
Met	Gln	Lys	Phe	Thr 470	Phe	Arg	Pro	Pro	Asn 475	Asn	Glu	Lys	Leu	Ser 480
Leu	Lys	Phe	Arg	Met 485	Gly	Ile	Thr	Ile	Ser 490	Pro	Val	Ser	His	Arg 495
Leu	Cys	Ala	Val	Pro	Gln	Val								

<210> 37 <211> 1577

<212> DNA

<213> Homo sapien

<400> 37

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<210> 38

<211> 338

<212> PRT

<213> Homo sapien

<400> 38

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His Leu Gly Arg Pro Ser Ala Gly Ala Val Val Ala His Pro Thr 20 25 30

Ser Gly Thr Ile Ser Ser Ala Ser Phe His Pro Gln Gln Phe Gln 35 40 45

Tyr Thr Leu Asp Asn Asn Val Leu Thr Leu Glu Gln Arg Lys Phe 50  $\phantom{0}55$   $\phantom{0}60$ 

Tyr Glu Glu Asn Gly Phe Leu Val Ile Lys Asn Leu Val Pro Asp
65 70 75

Ala Asp Ile Gln Arg Phe Arg Asn Glu Phe Glu Lys Ile Cys Arg 80 85 90

Lys Glu Val Lys Pro Leu Gly Leu Thr Val Met Arg Asp Val Thr 95 100 105

Ile Ser Lys Ser Glu Tyr Ala Pro Ser Glu Lys Met Ile Thr Lys 110 115 120

Val Gln Asp Phe Gln Glu Asp Lys Glu Leu Phe Arg Tyr Cys Thr 125 130 130

Leu Pro Glu Ile Leu Lys Tyr Val Glu Cys Phe Thr Gly Pro Asn 140 145 150

Ile Met Ala Met His Thr Met Leu Ile Asn Lys Pro Pro Asp Ser 155 160 165

Gly Lys Lys Thr Ser Arg His Pro Leu His Gln Asp Leu His Tyr 170 175 180

Phe Pro Phe Arg Pro Ser Asp Leu Ile Val Cys Ala Trp Thr Ala 185 190 195

Met Glu His Ile Ser Arg Asn Asn Gly Cys Leu Val Val Leu Pro 200 205 210

Gly Thr His Lys Gly Ser Leu Lys Pro His Asp Tyr Pro Lys Trp 215 220 225

Glu Gly Gly Val Asn Lys Met Phe His Gly Ile Gln Asp Tyr Glu 230 235 240 Glu Asn Lys Ala Arg Val His Leu Val Met Glu Lys Gly Asp Thr Val Phe Phe His Pro Leu Leu Ile His Gly Ser Gly Gln Asn Lys 265 270 Thr Gln Gly Phe Arg Lys Ala Ile Ser Cys His Phe Ala Ser Ala Asp Cys His Tyr Ile Asp Val Lys Gly Thr Ser Gln Glu Asn Ile 300 Glu Lys Glu Val Val Gly Ile Ala His Lys Phe Phe Gly Ala Glu Asn Ser Val Asn Leu Lys Asp Ile Trp Met Phe Arg Ala Arg Leu 325 Val Lys Gly Glu Arg Thr Asn Leu

<210> 39

<211> 716

<212> DNA

<213> Homo sapien

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<210> 40
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<211> 98

<212> PRT

<213> Homo sapien

## <400> 40

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Glu Ile Gln Lys His Asn His Ser Lys Ser Thr Trp Leu Ile Leu 20 25 30

His His Lys Val Tyr Asp Leu Thr Lys Phe Leu Glu Glu His Pro 35 40 45

Gly Glu Glu Val Leu Arg Glu Gln Ala Gly Gly Asp Ala Thr 50 55 60

Glu Asn Phe Glu Asp Val Gly His Ser Thr Asp Ala Arg Glu Met
65 70 75

Ser Lys Thr Phe Ile Ile Gly Glu Leu His Pro Asp Asp Arg Pro 80 85 90

Lys Leu Asn Lys Pro Pro Glu Pro

<210> 41

<211> 578

<212> DNA

<213> Homo sapien

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tetttgetga gggteacatt gagetgeagg ttgaateegg ggtgeettta 150
ggatteagea ceatggegga agaeatggag accaaaatea agaaetacaa 200
gaeegeeeet tttgaeagee getteeceaa eeagaaeeag aetagaaaet 250
getggeagaa etaeetggae tteeaeeget gteagaagge aatgaeeget 300
aaaggaggeg atatetetgt gtgegaatgg taeeagegtg tgtaeeagte 350
eetetgeeee acateetggg teaeagaetg ggatgageaa egggetgaag 400
geaegtttee egggaagate tgaaetgget geateteeet tteetetgte 450
eteeateett eteeeaggat ggtgaagggg gaeetggtae eeagtgatee 500
ceaeeeeagg ateetaaate atgaettaee tgetaataaa aaeteattgg 550
aaaagtgaaa aaaaaaaaa aaaaaaaa 578

<210> 42

<211> 86

<212> PRT

<213> Homo sapien

<210> 43 <211> 2444 <212> DNA

<213> Homo sapien

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<210> 44

<211> 596 <212> PRT <213> Homo sapien <400> 44 Met Pro Gln Leu Asn Gly Gly Gly Gly Asp Asp Leu Gly Ala Asn Asp Glu Leu Ile Ser Phe Lys Asp Glu Gly Glu Glu Glu Lys Ser Ser Glu Asn Ser Ser Ala Glu Arg Asp Leu Ala Asp Val Lys Ser Ser Leu Val Asn Glu Ser Glu Thr Asn Gln Asn Ser Ser Ser Asp Ser Glu Ala Glu Arg Arg Pro Pro Pro Arg Ser Glu Ser Phe Arg Asp Lys Ser Arg Glu Ser Leu Glu Glu Ala Ala Lys Arg Gln Asp Gly Gly Leu Phe Lys Gly Pro Pro Tyr Pro Gly Tyr Pro Phe Ile Met Ile Pro Asp Leu Thr Ser Pro Tyr Leu Pro Asn Gly Ser 115 Leu Ser Pro Thr Ala Arg Thr Tyr Leu Gln Met Lys Trp Pro Leu 125 130 135 Leu Asp Val Gln Ala Gly Ser Leu Gln Ser Arg Gln Ala Leu Lys 145 Asp Ala Arg Ser Pro Ser Pro Ala His Ile Val Ser Asn Lys Val 160 165 Pro Val Val Gln His Pro His His Val His Pro Leu Thr Pro Leu 170 175 Ile Thr Tyr Ser Asn Glu His Phe Thr Pro Gly Asn Pro Pro 185 190 195 His Leu Pro Ala Asp Val Asp Pro Lys Thr Gly Ile Pro Arg Pro 205 Pro His Pro Pro Asp Ile Ser Pro Tyr Tyr Pro Leu Ser Pro Gly 215 Thr Val Gly Gln Ile Pro His Pro Leu Gly Trp Leu Val Pro Gln 230 Gln Gly Gln Pro Val Tyr Pro Ile Thr Thr Gly Gly Phe Arg His 245 Pro Tyr Pro Thr Ala Leu Thr Val Asn Ala Ser Val Ser Arg Phe 260 265

Pro	Pro	His	Met	Val 275	Pro	Pro	His	His	Thr 280	Leu	His	Thr	Thr	Gly 285
Ile	Pro	His	Pro	Ala 290	Ile	Val	Thr	Pro	Thr 295	Val	Lys	Gln	Glu	Ser 300
Ser	Gln	Ser	Asp	Val 305	Gly	Ser	Leu	His	Ser 310	Ser	Lys	His	Gln	Asp 315
Ser	Lys	Lys	Glu	Glu 320	Glu	Lys	Lys	Lys	Pro 325	His	Ile	Lys	Lys	Pro 330
Leu	Asn	Ala	Phe	Met 335	Leu	Tyr	Met	Lys	Glu 340	Met	Arg	Ala	Lys	Val 345
Val	Ala	Glu	Cys	Thr 350	Leu	Lys	Glu	Ser	Ala 355	Ala	Ile	Asn	Gln	Ile 360
Leu	Gly	Arg	Arg	Trp 365	His	Ala	Leu	Ser	Arg 370	Glu	Glu	Gln	Ala	Lys 375
Tyr	Tyr	Glu	Leu	Ala 380	Arg	Lys	Glu	Arg	Gln 385	Leu	His	Met	Gln	Leu 390
Tyr	Pro	Gly	Trp	Ser 395	Ala	Arg	Asp	Asn	Tyr 400	Gly	Lys	Lys	Lys	Lys 405
Arg	Lys	Arg	Asp	Lys 410	Gln	Pro	Gly	Glu	Thr 415	Asn	Glu	His	Ser	Glu 420
Cys	Phe	Leu	Asn	Pro 425	Cys	Leu	Ser	Leu	Pro 430	Pro	Ile	Thr	Asp	Leu 435
Ser	Ala	Pro	Lys	Lys 440	Cys	Arg	Ala	Arg	Phe 445	Gly	Leu	Asp	Gln	Gln 450
Asn	Asn	Trp	Cys	Gly 455	Pro	Cys	Arg	Arg	Lys 460	Lys	Lys	Cys	Val	Arg 465
Tyr	Ile	Gln	Gly	Glu 470	Gly	Ser	Cys	Leu	Ser 475	Pro	Pro	Ser	Ser	Asp 480
Gly	Ser	Leu	Leu	Asp 485	Ser	Pro	Pro	Pro	Ser 490	Pro	Asn	Leu	Leu	Gly 495
Ser	Pro	Pro	Arg	Asp 500	Ala	Lys	Ser	Gln	Thr 505	Glu	Gln	Thr	Gln	Pro 510
Leu	Ser	Leu	Ser	Leu 515	Lys	Pro	Asp	Pro	Leu 520	Ala	His	Leu	Ser	Met 525
Met	Pro	Pro	Pro	Pro 530	Ala	Leu	Leu	Leu	Ala 535	Glu	Ala	Thr	His	Lys 540
Ala	Ser	Ala	Leu	Cys 545	Pro	Asn	Gly	Ala	Leu 550	Asp	Leu	Pro	Pro	Ala 555
Ala	Leu	Gln	Pro	Ala 560	Ala	Pro	Ser	Ser	Ser 565	Ile	Ala	Gln	Pro	Ser 570

Thr Ser Trp Leu His Ser His Ser Ser Leu Ala Gly Thr Gln Pro 575 580 585

Gln Pro Leu Ser Leu Val Thr Lys Ser Leu Glu 590 595

<210> 45

<211> 3697

<212> DNA

<213> Homo sapien

<400> 45

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t: --...

. . .

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<21Ü> 46

<211> 832

<212> PRT

<213> Homo sapien

## <400> 46

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Leu Lys Pro Met Thr Phe Ser Ile Tyr Glu Gly Gln Glu Pro Ser 35 40 45

Gln Ile Ile Phe Gln Phe Lys Ala Asn Pro Pro Ala Val Thr Phe
50 55 60

Glu Leu Thr Gly Glu Thr Asp Asn Ile Phe Val Ile Glu Arg Glu
65 70 75

Gly Leu Leu Tyr Tyr Asn Arg Ala Leu Asp Arg Glu Thr Arg Ser 80 85 90

Thr	His	Asn	Leu	Gln 95	Val	Ala	Ala	Leu	Asp 100	Ala	Asn	Gly	Ile	Ile 105
Val	Glu	Gly	Pro	Val 110	Pro	Ile	Thr	Ile	Glu 115	Val	Lys	Asp	Ile	Asn 120
Asp	Asn	Arg	Pro	Thr 125	Phe	Leu	Gln	Ser	Lys 130	Tyr	Glu	Gly	Ser	Val 135
Arg	Gln	Asn	Ser	Arg 140	Pro	Gly	Lys	Pro	Phe 145	Leu	Tyr	Val	Asn	Ala 150
Thr	Asp	Leu	Asp	Asp 155	Pro	Ala	Thr	Pro	Asn 160	Gly	Gln	Leu	Tyr	Tyr 165
Gln	Ile	Val	Ile	Gln 170	Leu	Pro	Met	Ile	Asn 175	Asn	Val	Met	Tyr	Phe 180
Gln	Ile	Asn	Asn	Lys 185	Thr	Gly	Ala	Ile	Ser 190	Leu	Thr	Arg	Glu	Gly 195
Ser	Gln	Glu	Leu	Asn 200	Pro	Ala	Lys	Asn	Pro 205	Ser	Tyr	Asn	Leu	Val 210
Ile	Ser	Val	Lys	Asp 215	Met	Gly	Gly	Gln	Ser 220	Glu	Asn	Ser	Phe	Ser 225
Asp	Thr	Thr	Ser	Val 230	Asp	Ile	Ile	Val	Thr 235	Glu	Asn	Ile	Trp	Lys 240
Ala	Pro	Lys	Pro	Val 245	Glu	Met	Val	Glu	Asn 250	Ser	Thr	Asp	Pro	His 255
Pro	Ile	Lys	Ile	Thr 260	Gln	Val	Arg	Trp	Asn 265	Asp	Pro	Gly	Ala	Gln 270
Tyr	Ser	Leu	Val	Asp 275	Lys	Glu	Lys	Leu	Pro 280	Arg	Phe	Pro	Phe	Ser 285
Ile	Asp	Gln	Glu	Gly 290	Asp	Ile	Tyr	Val	Thr 295	Gln	Pro	Leu	Asp	Arg 300
Glu	Glu	Lys	Asp	Ala 305	Tyr	Val	Phe	Tyr	Ala 310	Val	Ala	Lys	Asp	Glu 315
Tyr	Gly	Lys	Pro	Leu 320	Ser	Tyr	Pro	Leu	Glu 325	Ile	His	Val	Lys	Val 330
Lys	Asp	Ile	Asn	Asp 335	Asn	Pro	Pro	Thr	Cys 340	Pro	Ser	Pro	Val	Thr 345
Val	Phe	Glu	Val	Gln 350	Glu	Asn	Glu	Arg	Leu 355	Gly	Asn	Ser	Ile	Gly 360
Thr	Leu	Thr	Ala	His 365	Asp	Arg	Asp	Glu	Glu 370	Asn	Thr	Ala	Asn	Ser 375
Phe	Leu	Asn	Tyr	Arg 380	Ile	Val	Glu	Gln	Thr 385	Pro	Lys	Leu	Pro	Met 390

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Asp	Gly	Leu	Phe	Leu 395	Ile	Gln	Thr	Tyr	Ala 400	Gly	Met	Leu	Gln	Leu 405
Ala	Lys	Gln	Ser	Leu 410	Lys	Lys	Gln	Asp	Thr 415	Pro	Gln	Tyr	Asn	Leu 420
Thr	Ile	Glu	Val	Ser 425	Asp	Lys	Asp	Phe	Lys 430	Thr	Leu	Cys	Phe	Val 435
Gln	Ile	Asn	Val	Ile 440	Asp	Ile	Asn	Asp	Gln 445	Ile	Pro	Ile	Phe	Glu 450
Lys	Ser	Asp	Tyr	Gly 455	Asn	Leu	Thr	Leu	Ala 460	Glu	Asp	Thr	Asn	Ile 465
Gly	Ser	Thr	Ile	Leu 470	Thr	Ile	Gln	Ala	Thr 475	Asp	Ala	Asp	Glu	Pro 480
Phe	Thr	Gly	Ser	Ser 485	Lys	Ile	Leu	Tyr	His 490	Ile	Ile	Lys	Gly	Asp 495
Ser	Glu	Gly	Arg	Leu 500	Gly	Val	Asp	Thr	Asp 505	Pro	His	Thr	Asn	Thr 510
Gly	Tyr	Val	Ile	Ile 515	Lys	Lys	Pro	Leu	Asp 520	Phe	Glu	Thr	Ala	Ala 525
Val	Ser	Asn	Ile	Val 530	Phe	Lys	Ala	Glu	Asn 535	Pro	Glu	Pro	Leu	Val 540
Phe	Gly	Val	Lys	Tyr 545	Asn	Ala	Ser	Ser	Phe 550	Ala	Lys	Phe	Thr	Leu 555
Ile	Val	Thr	Asp	Val 560	Asn	Glu	Ala	Pro	Gln 565	Phe	Ser	Gln	His	Val 570
Phe	Gln	Ala	Lys	Val 575	Ser	Glu	Asp	Val	Ala 580	Ile	Gly	Thr	Lys	Val 585
Gly	Asn	Val	Thr	Ala 590	Lys	Asp	Pro	Glu	Gly 595	Leu	Asp	Ile	Ser	Tyr 600
Ser	Leu	Arg	Gly	Asp 605	Thr	Arg	Gly	Trp	Leu 610	Lys	Ile	Asp	His	Val 615
Thr	Gly	Glu	Ile	Phe 620	Ser	Val	Ala	Pro	Leu 625	Asp	Arg	Glu	Ala	Gly 630
Ser	Pro	Tyr	Arg	Val 635	Gln	Val	Val	Ala	Thr 640	Glu	Val	Gly	Gly	Ser 645
Ser	Leu	Ser	Ser	Val 650	Ser	Glu	Phe	His	Leu 655	Ile	Leu	Met	Asp	Val 660
Asn	Asp	Asn	Pro	Pro 665	Arg	Leu	Ala	Lys	Asp 670	Tyr	Thr	Gly	Leu	Phe 675
Phe	Cys	His	Pro	Leu 680	Ser	Ala	Pro	Gly	Ser 685	Leu	Ile	Phe	Glu	Ala 690
Thr	Asp	Asp	Asp	Gln	His	Leu	Phe	Arg	Gly	Pro	His	Phe	Thr	Phe

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<400> 48

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Thr Gly Cys Cys Ala Gly Gly Ala Ala Gly Cys Ala Gly Ala Gly
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Thr Gly Gly Ala Gly Ala Cys Gly Gly Ala Cys Cys Thr Gly Ala 95 100 105

Gly Cys Cys Cys Gly Ala Gly Gly Ala Ala Gly Ala Gly Cys  $110 \\ \hspace{1.5cm}115 \\ \hspace{1.5cm}120$ 

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Cys Thr Gly Ala Thr Thr Cys Cys Ala Cys Cys Cys Cys Ala Gly 140 145 150

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Суѕ	Cys	Ala	Gly	Gly 590	Ala	Ala	Ala	Gly	Cys 595	Cys	Ala	Ala	Gly	Cys 600
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20 25 30

Arg Leu Glu Leu Ser Asp Ile Tyr Gln Ile Pro Ser Val Asp Ser

35 40 45 Ala Asp Asn Leu Ser Glu Lys Leu Glu Arg Glu Trp Asp Arg Glu Leu Ala Ser Lys Lys Asn Pro Lys Leu Ile Asn Ala Leu Arg Arg Cys Phe Phe Trp Arg Phe Met Phe Tyr Gly Ile Phe Leu Tyr Leu Gly Glu Val Thr Lys Ala Val Gln Pro Leu Leu Gly Arg Ile 100 Ile Ala Ser Tyr Asp Pro Asp Asn Lys Glu Glu Arg Ser Ile Ala Ile Tyr Leu Gly Ile Gly Leu Cys Leu Leu Phe Ile Val Arg Thr 130 Leu Leu His Pro Ala Ile Phe Gly Leu His His Ile Gly Met Gln Met Arg Ile Ala Met Phe Ser Leu Ile Tyr Lys Lys Thr Leu Lys Leu Ser Ser Arg Val Leu Asp Lys Ile Ser Ile Gly Gln Leu Val Ser Leu Leu Ser Asn Asn Leu Asn Lys Phe Asp Glu Gly Leu Ala Leu Ala His Phe Val Trp Ile Ala Pro Leu Gln Val Ala Leu Leu Met Gly Leu Ile Trp Glu Leu Leu Gln Ala Ser Ala Phe Cys Gly Leu Gly Phe Leu Ile Val Leu Ala Leu Phe Gln Ala Gly Leu 230 Gly Arg Met Met Lys Tyr Arg Asp Gln Arg Ala Gly Lys Ile 250 Ser Glu Arg Leu Val Ile Thr Ser Glu Met Ile Glu Asn Ile Gln 260 265 Ser Val Lys Ala Tyr Cys Trp Glu Glu Ala Met Glu Lys Met Ile 280 Glu Asn Leu Arg Gln Thr Glu Leu Lys Leu Thr Arg Lys Ala Ala 290 295 · 300 Tyr Val Arg Tyr Phe Asn Ser Ser Ala Phe Phe Ser Gly Phe 305 310 Phe Val Val Phe Leu Ser Val Leu Pro Tyr Ala Leu Ile Lys Gly Ile Ile Leu Arg Lys Ile Phe Thr Thr Ile Ser Phe Cys Ile Val 340

200 1119	Met	Ala	Val 350	Thr	Arg	Gln	Phe	Pro 355	Trp	Ala	Val	Gln	Thr 360
Trp Tyr	Asp	Ser	Leu 365	Gly	Ala	Ile	Asn	Lys 370	Ile	Gln	Asp	Phe	Leu 375
Gln Lys	Gln	Glu	Tyr 380	Lys	Thr	Leu	Glu	Tyr 385	Asn	Leu	Thr	Thr	Thr 390
Glu Val	Val	Met	Glu 395	Asn	Val	Thr	Ala	Phe 400	Trp	Glu	Glu	Gly	Phe 405
Gly Glu	Leu	Phe	Glu 410	Lys	Ala	Lys	Gln	Asn 415	Asn	Asn	Asn	Arg	Lys 420
Thr Ser	Asn	Gly	Asp 425	Asp ·	Ser	Leu	Phe	Phe 430	Ser	Asn	Phe	Ser	Leu 435
Leu Gly	Thr	Pro	Val 440	Leu	Lys	Asp	Ile	Asn 445	Phe	Lys	Ile	Glu	Arg 450
Gly Gln	Leu	Leu	Ala 455	Val	Ala	Gly	Ser	Thr 460	Gly	Ala	Gly	Lys	Thr 465
Ser Leu	Leu	Met	Met 470	Ile	Met	Gly	Glu	Leu 475	Glu	Pro	Ser	Glu	Gly 480
Lys Ile	Lys	His	Ser 485	Gly	Arg	Ile	Ser	Phe 490	Cys	Ser	Gln	Phe	Ser 495
Trp Ile	Met	Pro	Gly 500	Thr	Ile	Lys	Glu	Asn 505	Ile	Ile	Phe	Gly	
			300					505					510
Ser Tyr	Asp	Glu		Arg	Tyr	Arg	Ser		Ile	Lys	Ala	Cys	
Ser Tyr			Tyr 515					Val 520				_	Gln 525
	Glu	Asp	Tyr 515 Ile 530	Ser	Lys	Phe	Ala	Val 520 Glu 535	Lys	Asp	Asn	Ile	Gln 525 Val 540
Leu Glu	Glu	Asp Gly	Tyr 515 Ile 530 Gly 545	Ser	Lys	Phe Leu	Ala Ser	Val 520 Glu 535 Gly 550	Lys Gly	Asp	Asn Arg	Ile	Gln 525 Val 540 Arg 555
Leu Glu	Glu Glu Leu	Asp Gly Ala	Tyr 515 Ile 530 Gly 545 Arg 560	Ser Ile Ala	Lys Thr	Phe Leu Tyr	Ala Ser Lys	Val 520 Glu 535 Gly 550 Asp 565	Lys Gly Ala	Asp Gln Asp	Asn Arg Leu	Ile Ala Tyr	Gln 525 Val 540 Arg 555 Leu 570
Leu Glu Leu Gly Ile Ser	Glu Glu Leu Ser	Asp Gly Ala Pro	Tyr 515 Ile 530 Gly 545 Arg 560 Phe 575	Ser Ile Ala	Lys Thr Val	Phe Leu Tyr Leu	Ala Ser Lys Asp	Val 520 Glu 535 Gly 550 Asp 565 Val	Lys Gly Ala Leu	Asp Gln Asp	Asn Arg Leu Glu	Ile Ala Tyr Lys	Gln 525 Val 540 Arg 555 Leu 570
Leu Glu Leu Gly Ile Ser Leu Asp	Glu Glu Leu Ser	Asp Gly Ala Pro	Tyr 515 Ile 530 Gly 545 Arg 560 Phe 575 Cys 590	Ser Ile Ala Gly Val	Lys Thr Val Tyr Cys	Phe Leu Tyr Leu Lys	Ala Ser Lys Asp	Val 520 Glu 535 Gly 550 Asp 565 Val 580 Met 595	Lys Gly Ala Leu Ala	Asp Gln Asp Thr	Asn Arg Leu Glu Lys	Ile Ala Tyr Lys Thr	Gln 525 Val 540 Arg 555 Leu 570 Glu 585 Arg 600
Leu Glu Leu Gly Ile Ser Leu Asp Ile Phe	Glu Glu Leu Ser Glu Val	Asp Gly Ala Pro Ser	Tyr 515 Ile 530 Gly 545 Arg 560 Phe 575 Cys 590 Ser 605	Ser Ile Ala Gly Val	Lys Thr Val Tyr Cys	Phe Leu Tyr Leu Lys	Ala Ser Lys Asp Leu	Val 520 Glu 535 Gly 550 Asp 565 Val 580 Met 595 Leu 610	Lys Gly Ala Leu Ala	Asp Gln Asp Thr Asn	Asn Arg Leu Glu Lys Ala	Ile Ala Tyr Lys Thr	Gln 525 Val 540 Arg 555 Leu 570 Glu 585 Arg 600 Lys

Gly	Cys	Asp	Ser	Phe 650	Asp	Gln	Phe	Ser	Ala 655	Glu	Arg	Arg	Asn	Ser 660
Ile	Leu	Thr	Glu	Thr 665	Leu	His	Arg	Phe	Ser 670	Leu	Glu	Gly	Asp	Ala 675
Pro	Val	Ser	Trp	Thr 680	Glu	Thr	Lys	Lys	Gln 685	Ser	Phe	Lys	Gln	Thr 690
Gly	Glu	Phe	Gly	Glu 695	Lys	Arg	Lys	Asn	Ser 700	Ile	Leu	Asn	Pro	Ile 705
Asn	Ser	Ile	Arg	Lys 710	Phe	Ser	Ile	Val	Gln 715	Lys	Thr	Pro	Leu	Gln 720
Met	Asn	Gly	Ile	Glu 725	Glu	Asp	Ser	Asp	Glu 730	Pro	Leu	Glu	Arg	Arg 735
Leu	Ser	Leu	Val	Pro 740	Asp	Ser	Glu	Gln	Gly 745	Glu	Ala	Ile	Leu	Pro 750
Arg	Ile	Ser	Val	Ile 755	Ser	Thr	Gly	Pro	Thr 760	Leu	Gln	Ala	Arg	Arg 765
Arg	Gln	Ser	Val	Leu 770	Asn	Leu	Met	Thr	His 775	Ser	Val	Asn	Gln	Gly 780
Gln	Asn	Ile	His	Arg 785	Lys	Thr	Thr	Ala	Ser 790	Thr	Arg	Lys	Val	Ser 795
Leu	Ala	Pro	Gln	Ala 800	Asn	Leu	Thr	Glu	Leu 805	Asp	Ile	Tyr	Ser	Arg 810
Arg	Leu	Ser	Gln	Glu 815	Thr	Gly	Leu	Glu	Ile 820	Ser	Glu	Glu	Ile	Asn 825
Glu	Glu	Asp	Leu	Lys 830	Glu	Cys	Leu	Phe	Asp 835	Asp	Met	Glu	Ser	Ile 840
Pro	Ala	Val	Thr	Thr 845	Trp	Asn	Thr	Tyr	Leu 850	Arg	Tyr	Ile	Thr	Val 855
His	Lys	Ser	Leu	Ile 860	Phe	Val	Leu	Ile	Trp 865	Cys	Leu	Val	Ile	Phe 870
Leu	Ala	Glu	Val	Ala 875	Ala	Ser	Leu	Val	Val 880	Leu	Trp	Leu	Leu	Gly 885
Asn	Thr	Pro	Leu	Gln 890	Asp	Lys	Gly	Asn	Ser 895	Thr	His	Ser	Arg	Asn 900
Asn	Ser	Tyr	Ala	Val 905	Ile	Ile	Thr	Ser	Thr 910	Ser	Ser	Tyr	Tyr	Val 915
Phe	Tyr	Ile	Tyr	Val 920	Gly	Val	Ala	Asp	Thr 925	Leu	Leu	Ala	Met	Gly 930
Phe	Phe	Arg	Gly	Leu 935	Pro	Leu	Val	His	Thr 940	Leu	Ile	Thr	Val	Ser 945
Lys	Ile	Leu	His	His	Lys	Met	Leu	His	Ser	Val	Leu	Gln	Ala	Pro

				950					955					960
Met	Ser	Thr	Leu	Asn	Thr	Leu	Lys	Ala	Gly	Gly	Ile	Leu	Asn	Arg

. - -

0 - 0

965 970 975

Phe Ser Lys Asp Ile Ala Ile Leu Asp Asp Leu Leu Pro Leu Thr 980 985 990

Ile Phe Asp Phe Ile Gln Leu Leu Ile Val Ile Gly Ala Ile 995 1000 1005

Ala Val Val Ala Val Leu Gln Pro Tyr Ile Phe Val Ala Thr Val
1010 1015 1020

Pro Val Ile Val Ala Phe Ile Met Leu Arg Ala Tyr Phe Leu Gln 1025 1030 1035

Thr Ser Gln Gln Leu Lys Gln Leu Glu Ser Glu Gly Arg Ser Pro 1040 1045 1050

Ile Phe Thr His Leu Val Thr Ser Leu Lys Gly Leu Trp Thr Leu 1055 1060 1065

Arg Ala Phe Gly Arg Gln Pro Tyr Phe Glu Thr Leu Phe His Lys 1070 1075 1080

Ala Leu Asn Leu His Thr Ala Asn Trp Phe Leu Tyr Leu Ser Thr 1085 1090 1095

Leu Arg Trp Phe Gln Met Arg Ile Glu Met Ile Phe Val Ile Phe 1100 1105 1110

Phe Ile Ala Val Thr Phe Ile Ser Ile Leu Thr Thr Gly Glu Gly 1115 1120 1125

Glu Gly Arg Val Gly Ile Ile Leu Thr Leu Ala Met Asn Ile Met
1130 1135 1140

Ser Thr Leu Gin Trp Ala Val Asn Ser Ser Ile Asp Val Asp Ser 1145 1150 1155

Leu Met Arg Ser Val Ser Arg Val Phe Lys Phe Ile Asp Met Pro 1160 1165 1170

Thr Glu Gly Lys Pro Thr Lys Ser Thr Lys Pro Tyr Lys Asn Gly 1175 1180 1185

Gln Leu Ser Lys Val Met Ile Ile Glu As<br/>n Ser His Val Lys Lys 1190 1195 1200

Asp Asp Ile Trp Pro Ser Gly Gly Gln Met Thr Val Lys Asp Leu 1205 1210 1215

Thr Ala Lys Tyr Thr Glu Gly Gly Asn Ala Ile Leu Glu Asn Ile 1220 1225 1230

Ser Phe Ser Ile Ser Pro Gly Gln Arg Val Gly Leu Leu Gly Arg 1235 1240 1245

Thr Gly Ser Gly Lys Ser Thr Leu Leu Ser Ala Phe Leu Arg Leu 1250 1255 1260 Leu Asn Thr Glu Gly Glu Ile Gln Ile Asp Gly Val Ser Trp Asp 1265 1270 1275

Ser Ile Thr Leu Gln Gln Trp Arg Lys Ala Phe Gly Val Ile Pro \$1280\$ \$1285\$ \$1290

Gln Lys Val Phe Ile Phe Ser Gly Thr Phe Arg Lys Asn Leu Asp 1295 1300 1305

Pro Tyr Glu Gln Trp Ser Asp Gln Glu Ile Trp Lys Val Ala Asp 1310 1315 1320

Glu Val Gly Leu Arg Ser Val Ile Glu Gln Phe Pro Gly Lys Leu 1325 1330 1335

Asp Phe Val Leu Val Asp Gly Gly Cys Val Leu Ser His Gly His 1340 1345 1350

Lys Gln Leu Met Cys Leu Ala Arg Ser Val Leu Ser Lys Ala Lys 1355 1360 1365

Ile Leu Leu Leu Asp Glu Pro Ser Ala His Leu Asp Pro Val Thr
1370 1375 1380

Tyr Gln Ile Ile Arg Arg Thr Leu Lys Gln Ala Phe Ala Asp Cys 1385 1390 1395

Thr Val Ile Leu Cys Glu His Arg Ile Glu Ala Met Leu Glu Cys 1400 1405 1410

Gln Gln Phe Leu Val Ile Glu Glu Asn Lys Val Arg Gln Tyr Asp 1415 1420 1425

Ser Ile Gln Lys Leu Leu Asn Glu Arg Ser Leu Phe Arg Gln Ala 1430 1435 1440

Ile Ser Pro Ser Asp Arg Val Lys Leu Phe Pro His Arg Asn Ser 1445 1450 1455

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<211> 1847

<212> DNA

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<400> 51

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<210> 52

<211> 359

<212> PRT

<213> Homo sapien

<400> 52

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Ala	Cys	Lys	Ile	Thr 20	Ile	Thr	Val		Leu 25	Ala	Val	Leu	Ile	Leu 30
Ile	Thr	Val	Ala	Gly 35	Asn	Val	Val	Val	Cys 40	Leu	Ala	Val	Gly	Leu 45

Asn Arg Arg Leu Arg Asn Leu Thr Asn Cys Phe Ile Val Ser Leu 50 55 60

Ala Ile Thr Asp Leu Leu Leu Gly Leu Leu Val Leu Pro Phe Ser 65 70 . 75

Ala Ile Tyr Gln Leu Ser Cys Lys Trp Ser Phe Gly Lys Val Phe  $80 \\ 85 \\ 90$ 

Cys Asn Ile Tyr Thr Ser Leu Asp Val Met Leu Cys Thr Ala Ser 95 100 105

Ile Leu Asn Leu Phe Met Ile Ser Leu Asp Arg Tyr Cys Ala Val110 115 120

Met Asp Pro Leu Arg Tyr Pro Val Leu Val Thr Pro Val Arg Val 125 130 130

Ala Ile Ser Leu Val Leu Ile Trp Val Ile Ser Ile Thr Leu Ser 140 145 150

Phe Leu Ser Ile His Leu Gly Trp Asn Ser Arg Asn Glu Thr Ser 155 160 165

Lys Gly Asn His Thr Thr Ser Lys Cys Lys Val Gln Val Asn Glu 170 175 180

Val Tyr Gly Leu Val Asp Gly Leu Val Thr Phe Tyr Leu Pro Leu 185 190 195

Leu Ile Met Cys Ile Thr Tyr Tyr Arg Ile Phe Lys Val Ala Arg 200 205 210

Asp Gln Ala Lys Arg Ile Asn His Ile Ser Ser Trp Lys Ala Ala 215 220 225

Thr Ile Arg Glu His Lys Ala Thr Val Thr Leu Ala Ala Val Met 230 235 240

Tyr Arg Gly Leu Arg Gly Asp Asp Ala Ile Asn Glu Val Leu Glu 260 265 Ala Ile Val Leu Trp Leu Gly Tyr Ala Asn Ser Ala Leu Asn Pro 275 280 285 Ile Leu Tyr Ala Ala Leu Asn Arg Asp Phe Arg Thr Gly Tyr Gln 295 Gln Leu Phe Cys Cys Arg Leu Ala Asn Arg Asn Ser His Lys Thr 310 315 Ser Leu Arg Ser Asn Ala Ser Gln Leu Ser Arg Thr Gln Ser Arg 320 Glu Pro Arg Gln Glu Glu Lys Pro Leu Lys Leu Gln Val Trp 340 345 Ser Gly Thr Glu Val Thr Ala Pro Gln Gly Ala Thr Asp Arg 350 355

<210> 53

<211> 5512

<212> DNA

<213> Homo sapien

<400> 53

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Asp His Phe Leu Ser Leu Gln Arg Met Phe Asn Asn Cys Glu Val
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Val Leu Gly Asn Leu Glu Ile Thr Tyr Val Gln Arg Asn Tyr Asp  $\phantom{000}65\phantom{000}70\phantom{000}75\phantom{000}$ 

Leu Ser Phe Leu Lys Thr Ile Gln Glu Val Ala Gly Tyr Val Leu  $80 \hspace{1cm} 85 \hspace{1cm} 90$ 

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Pro Met Arg Asn Leu Gln Glu Ile Leu His Gly Ala Val Arg Phe 140 145 150

Ser Asn Asn Pro Ala Leu Cys Asn Val Glu Ser Ile Gln Trp Arg 155 160 165

Asp Ile Val Ser Ser Asp Phe Leu Ser Asn Met Ser Met Asp Phe 170 175 180

Gln Asn His Leu Gly Ser Cys Gln Lys Cys Asp Pro Ser Cys Pro 185 190 195

Asn Gly Ser Cys Trp Gly Ala Gly Glu Glu Asn Cys Gln Lys Leu 200 205 210

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<210> 56

<211> 987

<212> PRT

<213> Homo sapien

<400> 56

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20 25 30

Ala Glu Leu Gly Trp Met Val His Pro Pro Ser Gly Trp Glu Glu
35 40 45

Val Ser Gly Tyr Asp Glu Asn Met Asn Thr Ile Arg Thr Tyr Gln 50 55 60

Val Cys Asn Val Phe Glu Ser Ser Gln Asn Asn Trp Leu Arg Thr  $\phantom{-}65\phantom{+}70\phantom{+}75\phantom{+}$ 

Lys Phe Ile Arg Arg Gly Ala His Arg Ile His Val Glu Met  $80 \hspace{1cm} 85 \hspace{1cm} 90$ 

Lys Phe Ser Val Arg Asp Cys Ser Ser Ile Pro Ser Val Pro Gly 95 100 105

Ser Cys Lys Glu Thr Phe Asn Leu Tyr Tyr Tyr Glu Ala Asp Phe 110 115 120

Asp Ser Ala Thr Lys Thr Phe Pro Asn Trp Met Glu Asn Pro Trp 125 130 135

Val Lys Val Asp Thr Ile Ala Ala Asp Glu Ser Phe Ser Gln Val

Asp Leu Gly Gly Arg Val Met Lys Ile Asn Thr Glu Val Arg Ser 155 160 165

Phe	Gly	Pro	Val	Ser 170	Arg	Ser	Gly	Phe	Tyr 175	Leu	Ala	Phe	Gln	Asp 180
Tyr	Gly	Gly	Cys	Met 185	Ser	Leu	Ile	Ala	Val 190	Arg	Val	Phe	Tyr	Arg 195
Lys	Cys	Pro	Arg	Ile 200	Ile	Gln	Asn	Gly	Ala 205	Ile	Phe	Gln	Glu	Thr 210
Leu	Ser	Gly	Ala	Glu 215	Ser	Thr	Ser	Leu	Val 220	Ala	Ala	Arg	Gly	Ser 225
Cys	Ile	Ala	Asn	Ala 230	Glu	Glu	Val	Asp	Val 235	Pro	Ile	Lys	Leu	Tyr 240
Cys	Asn	Gly	Asp	Gly 245	Glu	Trp	Leu	Val	Pro 250	Ile	Gly	Arg	Cys	Met 255
Cys	Lys	Ala	Gly	Phe 260	Glu	Ala	Val	Glu	Asn 265	Gly	Thr	Val	Cys	Arg 270
Gly	Cys	Pro	Ser	Gly 275	Thr	Phe	Lys	Ala	Asn 280	Gln	Gly	Asp	Glu	Ala 285
Cys	Thr	His	Cys	Pro 290	Ile	Asn	Ser	Arg	Thr 295	Thr	Ser	Glu	Gly	Ala 300
Thr	Asn	Cys	Val	Cys 305	Arg	Asn	Gly	Tyr	Tyr 310	Arg	Ala	Asp	Leu	Asp 315
Pro	Leu	Asp	Met	Pro 320	Cys	Thr	Thr	Ile	Pro 325	Ser	Ala	Pro	Gln	Ala 330
	Leu			320					325					330
Val		Ser	Ser	320 Val 335	Asn	Glu	Thr	Ser	325 Leu 340	Met	Leu	Glu	Trp	330 Thr 345
Val Pro	Ile	Ser	Ser Asp	320 Val 335 Ser 350	Asn	Glu Gly	Thr Arg	Ser Glu	325 Leu 340 Asp 355	Met Leu	Leu Val	Glu Tyr	Trp Asn	330 Thr 345 Ile 360
Val Pro Ile	Ile Pro	Ser Arg Lys	Ser Asp Ser	320 Val 335 Ser 350 Cys 365	Asn Gly Gly	Glu Gly Ser	Thr Arg Gly	Ser Glu Arg	325 Leu 340 Asp 355 Gly 370	Met Leu Ala	Leu Val Cys	Glu Tyr Thr	Trp Asn Arg	330 Thr 345 Ile 360 Cys 375
Val Pro Ile Gly	Ile Pro Cys	Ser Arg Lys Asn	Ser Asp Ser Val	320 Val 335 Ser 350 Cys 365 Gln 380	Asn Gly Gly Tyr	Glu Gly Ser Ala	Thr Arg Gly Pro	Ser Glu Arg	325 Leu 340 Asp 355 Gly 370 Gln 385	Met Leu Ala Leu	Leu Val Cys Gly	Glu Tyr Thr	Trp Asn Arg Thr	330 Thr 345 Ile 360 Cys 375 Glu 390
Val Pro Ile Gly Pro	Ile Pro Cys Asp	Ser Arg Lys Asn	Ser Asp Ser Val	320 Val 335 Ser 350 Cys 365 Gln 380 Ile 395	Asn Gly Gly Tyr	Glu Gly Ser Ala Asp	Thr Arg Gly Pro	Ser Glu Arg Arg	325 Leu 340 Asp 355 Gly 370 Gln 385 Ala 400	Met Leu Ala Leu His	Leu Val Cys Gly Thr	Glu Tyr Thr Leu Gln	Trp Asn Arg Thr	330 Thr 345 Ile 360 Cys 375 Glu 390 Thr 405
Val Pro Ile Gly Pro	Ile Pro Cys Asp	Ser Arg Lys Asn Ile	Ser Asp Ser Val Tyr Gln	320 Val 335 Ser 350 Cys 365 Gln 380 Ile 395 Ala 410	Asn Gly Gly Tyr Ser Val	Glu Gly Ser Ala Asp	Thr Arg Gly Pro Leu Gly	Ser Glu Arg Arg Leu Val	325 Leu 340 Asp 355 Gly 370 Gln 385 Ala 400 Thr 415	Met Leu Ala Leu His	Leu Val Cys Gly Thr	Glu Tyr Thr Leu Gln Ser	Trp Asn Arg Thr Tyr	330 Thr 345 Ile 360 Cys 375 Glu 390 Thr 405 Phe 420
Val Pro Ile Gly Pro Phe Ser	Ile Pro Cys Asp Arg	Ser Arg Lys Asn Ile Ile	Ser Asp Ser Val Tyr Gln Phe	320 Val 335 Ser 350 Cys 365 Gln 380 Ile 395 Ala 410 Ala 425	Asn Gly Gly Tyr Ser Val	Glu Gly Ser Ala Asp Asn Val	Thr Arg Gly Pro Leu Gly Asn	Ser Glu Arg Arg Leu Val	325 Leu 340 Asp 355 Gly 370 Gln 385 Ala 400 Thr 415 Thr 430	Met Leu Ala Leu His Asp	Leu Val Cys Gly Thr Gln	Glu Tyr Thr Leu Gln Ser	Trp Asn Arg Thr Tyr Pro	330 Thr 345 Ile 360 Cys 375 Glu 390 Thr 405 Phe 420 Ala 435

Ile	Leu	Asp	Tyr	Glu 470	Leu	Gln	Tyr	Tyr	Glu 475	Lys	Glu	Leu	Ser	Glu 480
Tyr	Asn	Ala	Thr	Ala 485	Ile	Lys	Ser	Pro	Thr 490	Asn	Thr	Val	Thr	Val 495
Gln	Gly	Leu	Lys	Ala 500	Gly	Ala	Ile	Tyr	Val 505	Phe	Gln	Val	Arg	Ala 510
Arg	Thr	Val	Ala	Gly 515	Tyr	Gly	Arg	Tyr	Ser 520	Gly	Lys	Met	Tyr	Phe 525
Gln	Thr	Met	Thr	Glu 530	Ala	Glu	Tyr	Gln	Thr 535	Ser	Ile	Gln	Glu	Lys 540
Leu	Pro	Leu	Ile	Ile 545	Gly	Ser	Ser	Ala	Ala 550	Gly	Leu	Val	Phe	Leu 555
Ile	Ala	Val	Val	Val 560	Ile	Ala	Ile	Val	Cys 565	Asn	Arg	Arg	Arg	Gly 570
Phe	Glu	Arg	Ala	Asp 575	Ser	Glu	Tyr	Thr	Asp 580	Lys	Leu	Gln	His	Tyr 585
Thr	Ser	Gly	His	Met 590	Thr	Pro	Gly	Met	Lys 595	Ile	Tyr	Ile	Asp	Pro 600
Phe	Thr	Tyr	Glu	Asp 605	Pro	Asn	Glu	Ala	Val 610	Arg	Glu	Phe	Ala	Lys 615
Glu	Ile	Asp	Ile	Ser 620	Cys	Val	Lys	Ile	Glu 625	Gln	Val	Ile	Gly	Ala 630
Gly	Glu	Phe	Gly	Glu 635	Val	Cys	Ser	Gly	His 640	Leu	Lys	Leu	Pro	Gly 645
Lys	Arg	Glu	Ile	Phe 650	Val	Ala	Ile	Lys	Thr 655	Leu	Lys	Ser	Gly	Tyr 660
Thr	Glu	Lys	Gln	Arg 665	Arg	Asp	Phe	Leu	Ser 670	Glu	Ala	Ser	Ile	Met 675
Gly	Gln	Phe	Asp	His 680	Pro	Asn	Val	Ile	His 685	Leu	Glu	Gly	Val	Val 690
Thr	Lys	Ser	Thr	Pro 695	Val	Met	Ile	Ile	Thr 700	Glu	Phe	Met	Glu	Asn 705
Gly	Ser	Leu	Asp	Ser 710	Phe	Leu	Arg	Gln	Asn 715	Asp	Gly	Gln	Phe	Thr 720
Val	Ile	Gln	Leu	Val 725	Gly	Met	Leu	Arg	Gly 730	Ile	Ala	Ala	Gly	Met 735
Lys	Tyr	Leu	Ala	Asp 740	Met	Asn	Tyr	Val	His 745	Arg	Asp	Leu	Ala	Ala 750
Arg	Asn	Ile	Leu	Val 755	Asn	Ser	Asn	Leu	Val 760	Cys	Lys	Val	Ser	Asp 765
Phe	Gly	.Leu	Ser	Arg	Phe	Leu	Glu	Asp	Asp	Thr	Ser	Asp	Pro	Thr

Tyr Thr Ser Ala Leu Gly Gly Lys Ile Pro Ile Arg Trp Thr Ala Pro Glu Ala Ile Gln Tyr Arg Lys Phe Thr Ser Ala Ser Asp Val Trp Ser Tyr Gly Ile Val Met Trp Glu Val Met Ser Tyr Gly Glu Arg Pro Tyr Trp Asp Met Thr Asn Gln Asp Val Ile Asn Ala Ile Glu Gln Asp Tyr Arg Leu Pro Pro Pro Met Asp Cys Pro Ser Ala Leu His Gln Leu Met Leu Asp Cys Trp Gln Lys Asp Arg Asn His Arg Pro Lys Phe Gly Gln Ile Val Asn Thr Leu Asp Lys Met Ile Arg Asn Pro Asn Ser Leu Lys Ala Met Ala Pro Leu Ser Ser Gly Ile Asn Leu Pro Leu Leu Asp Arg Thr Ile Pro Asp Tyr Thr Ser Phe Asn Thr Val Asp Glu Trp Leu Glu Ala Ile Lys Met Gly Gln Tyr Lys Glu Ser Phe Ala Asn Ala Gly Phe Thr Ser Phe Asp Val Val Ser Gln Met Met Met Glu Asp Ile Leu Arg Val Gly Val Thr Leu Ala Gly His Gln Lys Lys Ile Leu Asn Ser Ile Gln Val Met Arg Ala Gln Met Asn Gln Ile Gln Ser Val Glu Val 

<210> 57

<211> 2033

<212> DNA

<213> Homo sapien

<400> 57

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<210> 58

<211> 188

<212> PRT

<213> Homo sapien

<400> 58

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Ile Met Ala Ile Ser Lys Val Phe Glu Leu Gly Leu Val Ala Gly 20 25 30

Leu Gly His Gln Glu Phe Ala Arg Pro Ser Arg Gly Tyr Leu Ala
35 40 45

Phe Arg Asp Asp Ser Ile Trp Pro Gln Glu Glu Pro Ala Ile Arg
50 55 60

Pro Arg Ser Ser Gln Arg Val Pro Pro Met Gly Ile Gln His Ser
65 70 75

Lys Glu Leu Asn Arg Thr Cys Cys Leu Asn Gly Gly Thr Cys Met 80 85 90

Leu Gly Ser Phe Cys Ala Cys Pro Pro Ser Phe Tyr Gly Arg Asn 95 100 105

Cys Glu His Asp Val Arg Lys Glu Asn Cys Gly Ser Val Pro His 110 115 120

Asp Thr Trp Leu Pro Lys Lys Cys Ser Leu Cys Lys Cys Trp His 125 130 130

Gly Gln Leu Arg Cys Phe Pro Gln Ala Phe Leu Pro Gly Cys Asp 140 145 150

Gly Leu Val Met Asp Glu His Leu Val Ala Ser Arg Thr Pro Glu
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Leu Pro Pro Ser Ala Arg Thr Thr Thr Phe Met Leu Val Gly Ile 170 175 180

Cys Leu Ser Ile Gln Ser Tyr Tyr 185

<210> 59

<211> 3346

<212> DNA

<213> Homo sapien

<400> 59

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<210> 60

<211> 346

<212> PRT

<213> Homo sapien

<400> 60

Met Ala Arg Pro Gly Gln Arg Trp Leu Gly Lys Trp Leu Val Ala 1 5 10 15

Met Val Val Trp Ala Leu Cys Arg Leu Ala Thr Pro Leu Ala Lys
20 25 30

Asn Leu Glu Pro Val Ser Trp Ser Ser Leu Asn Pro Lys Phe Leu 35 40 45

Ser Gly Lys Gly Leu Val Ile Tyr Pro Lys Ile Gly Asp Lys Leu 50 55 60

Asp Ile Ile Cys Pro Arg Ala Glu Ala Gly Arg Pro Tyr Glu Tyr
65 70 75

Tyr Lys Leu Tyr Leu Val Arg Pro Glu Gln Ala Ala Cys Ser 80 85 90

Thr Val Leu Asp Pro Asn Val Leu Val Thr Cys Asn Arg Pro Glu 95 100 105

Gln Glu Ile Arg Phe Thr Ile Lys Phe Gln Glu Phe Ser Pro Asn 110 115 120

Tyr Met Gly Leu Glu Phe Lys Lys His His Asp Tyr Tyr Ile Thr 125 130 135

Ser Thr Ser Asn Gly Ser Leu Glu Gly Leu Glu Asn Arg Glu Gly 140 145 150

Gly Val Cys Arg Thr Arg Thr Met Lys Ile Ile Met Lys Val Gly
155 160 165

Gln Asp Pro Asn Ala Val Thr Pro Glu Gln Leu Thr Thr Ser Arg 170 175 180

Pro Ser Lys Glu Ala Asp Asn Thr Val Lys Met Ala Thr Gln Ala 185 190 , 195

Pro Gly Ser Arg Gly Ser Leu Gly Asp Ser Asp Gly Lys His Glu 200 205 210

Thr Val Asn Gln Glu Glu Lys Ser Gly Pro Gly Ala Ser Gly Gly

				215					220					225
Ser	Ser	Gly	Asp	Pro 230	Asp	Gly	Phe	Phe	Asn 235	Ser	Lys	Val	Ala	Leu 240
Phe	Ala	Ala	Val	Gly 245	Ala	Gly	Cys	Val	Ile 250	Phe	Leu	Leu	Ile	Ile 255
Ile	Phe	Leu	Thr	Val 260	Leu	Leu	Leu	Lys	Leu 265	Arg	Lys	Arg	His	Arg 270
Lys	His	Thr	Gln	Gln 275	Arg	Ala	Ala	Ala	Leu 280	Ser	Leu	Ser	Thr	Leu 285
Ala	Ser	Pro	Lys	Gly 290	Gly	Ser	Gly	Thr	Ala 295	Gly	Thr	Glu	Pro	Ser 300
Asp	Ile	Ile	Ile	Pro 305	Leu	Arg	Thr	Thr	Glu 310	Asn	Asn	Tyr	Cys	Pro 315
His	Tyr	Glu	Lys	Val 320	Ser	Gly	Asp	Tyr	Gly 325	His	Pro	Val	Tyr	Ile 330
Val	Gln	Glu	Met	Pro 335	Pro	Gln	Ser	Pro	Ala 340	Asn	Ile	Tyr	Tyr	Lys 345

Val

<210> 61 <211> 2438 <212> DNA

<213> Homo sapien

## <400> 61

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etetegeggt tgeegetget geegetgeeg etgetgetge tgetggeget 200
ggggaceege gggggetgeg eegegeeege aeeeggeeg egegeeggag 250
aceteageet gggagtggag tggetaagea ggtteggtta eetgeeeeg 300
getgaceea eaaeagggea getgeagaeg eaagaggage tgtetaagge 350
cateacagee atgeageat ttggtggeet ggaggeeaee ggeateetgg 400
aegaggeeae eetggeeetg atgaaaaeee eaegetgete eetgeeagae 450
eteeetgtee tgaeeeagge tegeaggaga egeeaggete eageeeeae 500
caagtggaae aagaggaaee tgtegtggag ggteeggaeg tteeeaegg 550
aeteaceaet ggggeaegae aeggtgegtg eaeteatgta etaegeete 600
aaggtetgga gegaeattge geeeetgaae tteeaegag tggeggeag 650

caccgccgac atccagatcg acttctccaa ggccgaccat aacgacggct 700 accepttega eggeeegge ggeaeegtgg eceaegeett etteeeegge 750 caccaccaca ccgccgggga cacccacttt gacgatgacg aggcctggac 800 etteegetee teggatgeee aegggatgga cetgtttgea gtggetgtee 850 acgagtttgg ccacgccatt gggttaagcc atgtggccgc tgcacactcc 900 atcatgcggc cgtactacca gggcccggtg ggtgacccgc tgcgctacgg 950 gctcccctac gaggacaagg tgcgcgtctg gcagctgtac ggtgtgcggg 1000 agtotgtgtc toccacggcg cagoocgagg agcotocoot gotgooggag 1050 ccccagaca accggtccag cgccccgccc aggaaggacg tgcccacag 1100 atgcagcact cactttgacg cggtggccca gatccgcggt gaagctttct 1150 tetteaaagg caagtaette tggeggetga egegggaeeg geaeetggtg 1200 tecetgeage eggeacagat geacegette tggeggggee tgeegetgea 1250 cctggacagc gtggacgccg tgtacgagcg caccagcgac cacaagatcg 1300 tcttctttaa aggagacagg tactgggtgt tcaaggacaa taacgtagag 1350 gaaggatacc cgcgccccgt ctccgacttc agcctcccgc ctggcggcat 1400 cgacgctgcc ttctcctggg cccacaatga caggacttat ttctttaagg 1450 accagetgta etggegetae gatgaceaea egaggeaeat ggaceeegge 1500 tacccegccc agagececct gtggaggggt gteeceagea egetggaega 1550 cgccatgcgc tggtccgacg gtgcctccta cttcttccgt ggccaggagt 1600 actggaaagt gctggatggc gagctggagg tggcacccgg gtacccacag 1650 tccacggccc gggactggct ggtgtgtgga gactcacagg ccgatggatc 1700 tgtggctgcg ggcgtggacg cggcagaggg gccccgcgcc cctccaggac 1750 aacatgacca gagccgctcg gaggacggtt acgaggtctg ctcatgcacc 1800 tetggggeat cetetecece gggggeeeca ggeeeactgg tggetgeeac 1850 catgctgctg ctgctgccgc cactgtcacc aggcgccctg tggacagcgg 1900 eccaggeeet gaegetatga cacacagege gageeeatga gaggacagag 1950 gcggtgggac agcctggcca cagagggcaa ggactgtgcc ggagtccctg 2000 ggggaggtgc tggcgcggga tgaggacggg ccaccetggc accggaaggc 2050 cagcagaggg cacggcccgc cagggctggg caggctcagg tggcaaggac 2100 ggagctgtcc cctagtgagg gactgtgttg actgacgagc cgaggggtgg 2150 ccgctccaga agggtgccca gtcaggccgc accgccgcca gcctcctccg 2200

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<210> 62

<211> 606

<212> PRT

<213> Homo sapien

<400> 62

Met Arg Arg Arg Ala Ala Arg Gly Pro Gly Pro Pro Pro Gly
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Pro Gly Leu Ser Arg Leu Pro Leu Leu Pro Leu Pro Leu Leu Leu 20 25 30

Leu Leu Ala Leu Gly Thr Arg Gly Gly Cys Ala Ala Pro Ala Pro 35 40 45

Ala Pro Arg Ala Glu Asp Leu Ser Leu Gly Val Glu Trp Leu Ser 50 55 60

Arg Phe Gly Tyr Leu Pro Pro Ala Asp Pro Thr Thr Gly Gln Leu 65 70 75

Gln Thr Gln Glu Glu Leu Ser Lys Ala Ile Thr Ala Met Gln Gln 80 85 90

Phe Gly Gly Leu Glu Ala Thr Gly Ile Leu Asp Glu Ala Thr Leu 95 100 105

Ala Leu Met Lys Thr Pro Arg Cys Ser Leu Pro Asp Leu Pro Val 110 115 120

. 15

Leu Thr Gln Ala Arg Arg Arg Gln Ala Pro Ala Pro Thr Lys 125 130 135

Trp Asn Lys Arg Asn Leu Ser Trp Arg Val Arg Thr Phe Pro Arg 140 145 150

Asp Ser Pro Leu Gly His Asp Thr Val Arg Ala Leu Met Tyr Tyr 155 160 165

Ala Leu Lys Val Trp Ser Asp Ile Ala Pro Leu Asn Phe His Glu 170 175 180

Val Ala Gly Ser Thr Ala Asp Ile Gln Ile Asp Phe Ser Lys Ala 185 190 195

Asp His Asn Asp Gly Tyr Pro Phe Asp Gly Pro Gly Gly Thr Val 200 205 210

Ala His Ala Phe Phe Pro Gly His His His Thr Ala Gly Asp Thr 215 220 225

His	Phe	Asp	Asp	Asp 230	Glu	Ala	Trp	Thr	Phe 235	Arg	Ser	Ser	Asp	Ala 240
His	Gly	Met	Asp	Leu 245	Phe	Ala	Val	Ala	Val 250	His	Glu	Phe	Gly	His 255
Ala	Ile	Gly	Leu	Ser 260	His	Val	Ala	Ala	Ala 265	His	Ser	Ile	Met	Arg 270
Pro	Tyr	Tyr	Gln	Gly 275	Pro	Val	Gly	Asp	Pro 280	Leu	Arg	Tyr	Gly	Leu 285
Pro	Tyr	Glu	Asp	Lys 290	Val	Arg	Val	Trp	Gln 295	Leu	Tyr	Gly	Val	Arg 300
Glu	Ser	Val	Ser	Pro 305	Thr	Ala	Gln	Pro	Glu 310	Glu	Pro	Pro	Leu	Leu 315
Pro	Glu	Pro	Pro	Asp 320	Asn	Arg	Ser	Ser	Ala 325	Pro	Pro	Arg	Lys	Asp 330
Val	Pro	His	Arg	Cys 335	Ser	Thr	His	Phe	Asp 340	Ala	Val	Ala	Gln	Ile 345
Arg	Gly	Glu	Ala	Phe 350	Phe	Phe	Lys	Gly	Lys 355	Tyr	Phe	Trp	Arg	Leu 360
Thr	Arg	Asp	Arg	His 365	Leu	Val	Ser	Leu	Gln 370	Pro	Ala	Gln	Met	His 375
Arg	Phe	Trp	Arg	Gly 380	Leu	Pro	Leu	His	Leu 385	Asp	Ser	Val	Asp	Ala 390
			Arg Arg	380					385					390
Val	Tyr	Glu		380 Thr 395	Ser	Asp	His	Lys	385 Ile 400	Val	Phe	Phe	Lys	390 Gly 405
Val Asp	Tyr Arg	Glu Tyr	Arg	380 Thr 395 Val 410	Ser Phe	Asp Lys	His Asp	Lys Asn	385 Ile 400 Asn 415	Val Val	Phe Glu	Phe Glu	Lys Gly	390 Gly 405 Tyr 420
Val Asp Pro	Tyr Arg Arg	Glu Tyr Pro	Arg Trp	380 Thr 395 Val 410 Ser 425	Ser Phe Asp	Asp Lys Phe	His Asp Ser	Lys Asn Leu	385 Ile 400 Asn 415 Pro 430	Val Val Pro	Phe Glu Gly	Phe Glu Gly	Lys Gly Ile	390 Gly 405 Tyr 420 Asp 435
Val Asp Pro	Tyr Arg Arg	Glu Tyr Pro	Arg Trp Val	380 Thr 395 Val 410 Ser 425 Trp 440	Ser Phe Asp	Asp Lys Phe	His Asp Ser Asn	Lys Asn Leu Asp	385 Ile 400 Asn 415 Pro 430 Arg 445	Val Val Pro	Phe Glu Gly Tyr	Phe Glu Gly Phe	Lys Gly Ile Phe	390 Gly 405 Tyr 420 Asp 435 Lys 450
Val Asp Pro Ala Asp	Tyr Arg Arg Ala Gln	Glu Tyr Pro Phe Leu	Arg Trp Val Ser	380 Thr 395 Val 410 Ser 425 Trp 440 Trp 455	Ser Phe Asp Ala	Asp Lys Phe His	His Asp Ser Asn	Lys Asn Leu Asp	385 Ile 400 Asn 415 Pro 430 Arg 445 His 460	Val Val Pro Thr	Phe Glu Gly Tyr Arg	Phe Glu Gly Phe His	Lys Gly Ile Phe Met	390 Gly 405 Tyr 420 Asp 435 Lys 450
Val Asp Pro Ala Asp	Tyr Arg Arg Ala Gln Gly	Glu Tyr Pro Phe Leu	Arg Trp Val Ser	380 Thr 395 Val 410 Ser 425 Trp 440 Trp 455 Ala 470	Ser Phe Asp Ala Arg	Asp Lys Phe His Tyr	His Asp Ser Asn Asp	Lys Asn Leu Asp Asp	385 Ile 400 Asn 415 Pro 430 Arg 445 His 460 Trp 475	Val Val Pro Thr Thr	Phe Glu Gly Tyr Arg	Phe Glu Gly Phe His	Lys Gly Ile Phe Met	390 Gly 405 Tyr 420 Asp 435 Lys 450 Asp 465 Ser 480
Val Asp Pro Ala Asp Pro	Tyr Arg Arg Ala Gln Gly Leu	Glu Tyr Pro Phe Leu Tyr Asp	Arg Trp Val Ser Tyr Pro	380 Thr 395 Val 410 Ser 425 Trp 440 Trp 455 Ala 470 Ala 485	Ser Phe Asp Ala Arg Gln Met	Asp Lys Phe His Tyr Ser	His Asp Ser Asn Asp Pro	Lys Asn Leu Asp Asp Leu Ser	385 Ile 400 Asn 415 Pro 430 Arg 445 His 460 Trp 475 Asp 490	Val Val Pro Thr Arg Gly	Phe Glu Gly Tyr Arg Gly Ala	Phe Glu Gly Phe His	Lys Gly Ile Phe Met Pro	390 Gly 405 Tyr 420 Asp 435 Lys 450 Asp 465 Ser 480 Phe 495

Cys Gly Asp Ser Gln Ala Asp Gly Ser Val Ala Ala Gly Val Asp 530 535 540 Ala Ala Glu Gly Pro Arg Ala Pro Pro Gly Gln His Asp Gln Ser 545 550 555 Arg Ser Glu Asp Gly Tyr Glu Val Cys Ser Cys Thr Ser Gly Ala 560 565 Ser Ser Pro Pro Gly Ala Pro Gly Pro Leu Val Ala Ala Thr Met 575 580 585 Leu Leu Leu Pro Pro Leu Ser Pro Gly Ala Leu Trp Thr Ala 595 600 Ala Gln Ala Leu Thr Leu

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<210> 63 <211> 1009 <212> DNA <213> Homo sapien

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aaacgctact gagtcacaat aaagattgtt ttaaagagta aaaaaaaaa 1000 aaaaaaaaa 1009

aaaaa	aaa	a 10	009											
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Phe A	la	Val	Pro	Val 20	Pro	Pro	Ala	Ala	Asp 25	His	Lys	Gly	Trp	Asp 30
Phe V	'al	Glu	Gly	Tyr 35	Phe	His	Gln	Phe	Phe 40	Leu	Thr	Glu	Lys	Glu 45
Ser P	ro	Leu	Leu	Thr 50	Gln	Glu	Thr	Gln	Thr 55	Gln	Leu	Leu	Gln	Gln 60
Phe H	is	Arg	Asn	Gly 65	Thr	Asp	Leu	Leu	Asp 70	Met	Gln	Met	His	Ala 75
Leu L	eu	His	Gln	Pro 80	His	Cys	Gly	Val	Pro 85	Asp	Gly	Ser	Asp	Thr 90
Ser I	le	Ser	Pro	Gly 95	Arg	Cys	Lys	Trp	Asn 100	Lys	His	Thr	Leu	Thr 105
Tyr A	rg	Ile	Ile	Asn 110	Tyr	Pro	His	Asp	Met 115	Lys	Pro	Ser	Ala	Val 120
Lys A	.sp	Ser	Ile	Tyr 125	Asn	Ala	Val	Ser	Ile 130	Trp	Ser	Asn	Val	Thr 135
Pro L	eu	Ile	Phe	Gln 140	Gln	Val	Gln	Asn	Gly 145	Asp	Ala	Asp	Ile	Lys 150
Val S	er	Phe	Trp	Gln 155	Trp	Ala	His	Glu	Asp 160	Gly	Trp	Pro	Phe	Asp 165
Gly P	ro	Gly	Gly	Ile 170	Leu	Gly	His		Phe 175	Leu	Pro	Asn	Ser	Gly 180
Asn P	ro	Gly	Val	Val 185	His	Phe	Asp	Lys	Asn 190	Glu	His	Trp	Ser	Ala 195
Ser A	sp	Thr	Gly	Tyr 200	Asn	Leu	Phe	Leu	Val 205	Ala	Thr	His	Glu	Ile 210
Gly H	is	Ser	Leu	Gly 215	Leu	Gln	His	Ser	Gly 220	Asn	Gln	Ser	Ser	Ile 225
Met T	yr	Pro	Thr	Tyr 230	Trp	Tyr	His	Asp	Pro 235	Arg	Thr	Phe	Gln	Leu 240
Ser A	la .	Asp	Asp	Ile 245	Gln	Arg	Ile	Gln	His 250	Leu	Tyr	Gly	Glu	Lys

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<210> 65 <211> 3410

<212> DNA

<213> Homo sapien

<400> 65

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<210> 66 <211> 748 <212> PRT <213> Homo sapien

## <400> 66

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Phe	His	Ser	Val	Ile 155	Tyr	His	Glu	Asp	Asp 160	Ile	Asn	Tyr	Pro	His 165
Lys	Tyr	Gly	Pro	Gln 170	Gly	Gly	Cys	Ala	Asp 175	His	Ser	Val	Phe	Glu 180
Arg	Met	Arg	Lys	Tyr 185	Gln	Met	Thr	Gly	Val 190	Glu	Glu	Val	Thr	Gln 195
Ile	Pro	Gln	Glu	Glu 200	His	Ala	Ala	Asn	Gly 205	Pro	Glu	Leu	Leu	Arg 210
Lys	Lys	Arg	Thr	Thr 215	Ser	Ala	Glu	Lys	Asn 220	Thr	Cys	Gln	Leu	Tyr 225
Ile	Gln	Thr	Asp	His 230	Leu	Phe	Phe	Lys	Tyr 235	Tyr	Gly	Thr	Arg	Glu 240
Ala	Val	Ile	Ala	Gln 245	Ile	Ser	Ser	His	Val 250	Lys	Ala	Ile	Asp	Thr 255
Ile	Tyr	Gln	Thr	Thr 260	Asp	Phe	Ser	Gly	Ile 265	Arg	Asn	Ile	Ser	Phe 270
Met	Val	Lys	Arg	Ile 275	Arg	Ile	Asn	Thr	Thr 280	Ala	Asp	Glu	Lys	Asp 285
Pro	Thr	Asn	Pro	Phe 290	Arg	Phe	Pro	Asn	Ile 295	Gly	Val	Glu	Lys	Phe 300
Leu	Glu	Leu	Asn	Ser 305	Glu	Gln	Asn	His	Asp 310	Asp	Tyr	Cys	Leu	Ala 315
Tyr	Val	Phe	Thr		Arg	Asp	Phe	Asp	Asp	Gly	Val	Leu	Gly	
				320					325					330
Ala	Trp	Val	Gly		Pro	Ser	Gly			Gly	Gly		Cys	
			Gly Leu	Ala 335			_	Ser	Ser 340	_		Ile	_	Glu 345
Lys	Ser	Lys		Ala 335 Tyr 350	Ser	Asp	Gly	Ser Lys	Ser 340 Lys 355	Lys	Ser	Ile Leu	Asn	Glu 345 Thr 360
Lys	Ser	Lys Ile	Leu	Ala 335 Tyr 350 Val 365	Ser Gln	Asp Asn	Gly Tyr	Ser Lys Gly	Ser 340 Lys 355 Ser 370	Lys His	Ser	Ile Leu Pro	Asn Pro	Glu 345 Thr 360 Lys 375
Lys Gly Val	Ser Ile Ser	Lys Ile His	Leu Thr	Ala 335 Tyr 350 Val 365 Thr 380	Ser Gln Phe	Asp Asn Ala	Gly Tyr His	Ser Lys Gly	Ser 340 Lys 355 Ser 370 Val 385	Lys His Gly	Ser Val His	Ile Leu Pro Asn	Asn Pro	Glu 345 Thr 360 Lys 375 Gly 390
Lys Gly Val Ser	Ser Ile Ser Pro	Lys Ile His	Leu Thr	Ala 335 Tyr 350 Val 365 Thr 380 Ser 395	Ser Gln Phe Gly	Asp Asn Ala Thr	Gly Tyr His	Ser Lys Gly Glu Cys	Ser 340 Lys 355 Ser 370 Val 385 Thr 400	Lys His Gly Pro	Ser Val His	Ile Leu Pro Asn Glu	Asn Pro Phe Ser	Glu 345 Thr 360 Lys 375 Gly 390 Lys 405
Lys Gly Val Ser Asn	Ser Ile Ser Pro	Lys Ile His Gly	Leu Thr Ile Asp	Ala 335 Tyr 350 Val 365 Thr 380 Ser 395 Lys 410	Ser Gln Phe Gly	Asp Asn Ala Thr	Gly Tyr His Glu	Ser Lys Gly Glu Cys Asn	Ser 340 Lys 355 Ser 370 Val 385 Thr 400	Lys His Gly Pro	Ser Val His Gly Met	Ile Leu Pro Asn Glu	Asn Pro Phe Ser Ala	Glu 345 Thr 360 Lys 375 Gly 390 Lys 405 Arg 420

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Asp	Glu	Cys	Cys	Phe 485	Asp	Ala	Asn	Gln	Pro 490	Glu	Gly	Arg	Lys	Cys 495
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Cys	Thr	Ala	Gln	Cys 515	Ala	Phe	Lys	Ser	Lys 520	Ser	Glu	Lys	Cys	Arg 525
Asp	Asp	Ser	Asp	Cys 530	Ala	Arg	Glu	Gly	Ile 535	Cys	Asn	Gly	Phe	Thr 540
Ala	Leu	Cys	Pro	Ala 545	Ser	Asp	Pro	Lys	Pro 550	Asn	Phe	Thr	Asp	Cys 555
Asn	Arg	His	Thr	Gln 560	Val	Cys	Ile	Asn	Gly 565	Gln	Cys	Ala	Gly	Ser 570
Ile	Cys	Glu	Lys	Tyr 575	Gly	Leu	Glu	Glu	Cys 580	Thr	Cys	Ala	Ser	Ser 585
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Lys	Met	Asp	Pro	Ser 605	Thr	Cys	Ala	Ser	Thr 610	Gly	Ser	Val	Gln	Trp 615
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Ile	Phe	Ser	Pro	Glu 665	Leu	Tyr	Glu	Asn	Ile 670	Ala	Glu	Trp	Ile	Val 675
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Leu	Met	Ala	Gly	Phe 695	Ile	Lys	Ile	Cys	Ser 700	Val	His	Thr	Pro	Ser 705
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Lys	Arg	Arg	Arg	Pro 725	Pro	Gln	Pro	Ile	Gln 730	Gln	Pro	Gln	Arg	Gln 735
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<sup>&</sup>lt;210> 68

<sup>&</sup>lt;211> 4498

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapien

<4000 Met 1		Met	Ala	Val 5	Gly	Asp	Arg	Ile	Tyr 10	Asp	Leu	Val	Met	Gln 15
Ile	Gly	Leu	Pro	Ala 20	Asp	Gly	Phe	Arg	Leu 25	Phe	Arg	Arg	Gly	Ala 30
Ala	Trp	Gln	Thr	Leu 35	Phe	Leu	Leu	Cys	Ala 40	Leu	Ala	Tyr	Cys	Ile 45
Asn	Glu	Ala	Ser	Ser 50	Glu	Gly	Arg	Val	Val 55	Cys	Tyr	Tyr	Thr	Asn 60
Trp	Ser	Val	Tyr	Arg 65	Pro	Gly	Thr	Ala	Lys 70	Phe	Asn	Pro	Gln	Asn 75
Ile	Asn	Pro	Tyr	Leu 80	Cys	Thr	His	Leu	Val 85	Tyr	Ala	Phe	Gly	Gly 90
Phe	Thr	Lys	Asp	Asn 95	Gln	Met	Lys	Pro	Phe 100	Asp	Lys	Tyr	Gln	Asp 105
Ile	Glu	Gln	Gly	Gly 110	Tyr	Ala	Lys	Phe	Thr 115	Gly	Leu	Ьуs	Thr	Tyr 120
Asn	Lys	Gln	Leu	Lys 125	Thr	Met	Ile	Ala	Ile 130	Gly	Gly	Trp	Asn	Glu 135
Ala	Ser	Ser	Arg	Phe 140	Ser	Pro	Leu	Val	Ala 145	Ser	Asn	Glu	Arg	Arg 150
Gln	Gln	Phe	Ile	Lys 155	Asn	Ile	Leu	Lys	Phe 160	Leu	Arg	Gln	Asn	His 165
Phe	Asp	Gly	Ile	Asp 170	Leu	Asp	Trp	Glu	Tyr 175	Pro	Ala	His	Arg	Glu 180
Gly	Gly.	Lys	Ser	Arg 185	Asp	Arg	Asp	Asn	Tyr 190	Ala	Gln	Phe	Val	Gln 195
Glu	Leu	Arg	Ala	Glu 200	Phe	Glu	Arg	Glu	Ala 205	Glu	Lys	Thr	Gly	Arg 210
Thr	Arg	Leu	Leu	Leu 215	Thr	Met	Ala	Val	Pro 220	Ala	Gly	Ile	Glu	Tyr 225
Ile	Asp	Lys	Gly	Tyr 230	Asp	Val	Pro	Lys	Leu 235	Asn	Lys	Tyr	Leu	Asp 240
Trp	Phe	Asn	Val	Leu 245	Thr	Tyr	Asp	Phe	His 250	Ser	Ser	His	Glu	Pro 255
Ser	Val	Asn	His	His 260	Ala	Pro	Leu	Tyr	Ser 265	Leu	Glu	Glu	Asp	Ser 270
Glu	Tyr	Asn	Tyr	Asp 275	Ala	Glu	Leu	Asn	Ile 280	Asp	Tyr	Ser	Ile	Lys 285
Tyr	Tyr	Leu	Lys	Ala 290	Gly	Ala	Asp	Arg	Asp 295	Lys	Leu	Val	Leu	Gly 300

Ile	Pro	Thr	Tyr	Gly 305	Arg	Ser	Tyr	Thr	Leu 310	Ile	Asn	Glu	Glu	Ser 315
Thr	Glu	Leu	Gly	Ala 320	Pro	Ala	Glu	Gly	Pro 325	Gly	Glu	Gln	Gly	Asp 330
Ala	Thr	Arg	Glu	Lys 335	Gly	Tyr	Leu	Ala	Tyr 340	Tyr	Glu	Ile	Cys	Gln 345
Thr	Leu	Lys	Asp	Asp 350	Pro	Glu	Trp	Thr	Val 355	Val	Gln	Pro	Asn	Ala 360
Asn	Val	Met	Gly	Pro 365	Tyr	Ala	Tyr	Arg	Arg 370	Asn	Gln	Trp	Val	Gly 375
Tyr	Asp	Asp	Glu	Ala 380	Ile	Val	Arg	Lys	Lys 385	Ala	Glu	Tyr	Val	Val 390
Ala	Gln	Gly	Leu	Gly 395	Gly	Ile	Met	Phe	Trp 400	Ala	Ile	Asp	Asn	Asp 405
Asp	Phe	Arg	Gly	Thr 410	Cys	Asn	Gly	Lys	Pro 415	Tyr	Pro	Leu	Ile	Glu 420
Ala	Ala	Lys	Glu	Ala 425	Met	Val	Glu	Ala	Leu 430	Gly	Leu	Gly	Ile	Asn 435
Glu	Val	Ala	Lys	Pro 440	Ser	Gly	Pro	Gln	Lys 445	Pro	Ser	Arg	Ser	Arg 450
Ser	Arg	Asp	Asn	Ala 455	Ser	Asn	Arg	Asn	Arg 460	Leu	Asn	Gly	Lys	Thr 465
Glu	Ala	Pro	Leu	Ser 470	Ser	Arg	Arg	Pro	Ser 475	Ala	Thr	Arg	Arg	Pro 480
Ala	Val	Ser	Ser	Thr 485	Gln	Ala	Pro	Pro	Pro 490	Ser	Thr	Thr	Phe	Lys 495
Leu	Thr	Glu	Ala	Glu 500	Gly	Ser	Ser	Leu	Tyr 505	Ile	Gly	Gly	Arg	Ala 510
Ser	Thr	Thr	Pro	Pro 515	Pro	Pro	Thr	Thr	Pro 520	Asp	Pro	Gly	Ser	Asp 525
Phe	Lys	Cys	Glu	Glu 530	Glu	Gly	Phe	Phe	Gln 535	His	Pro	Arg	Asp	Cys 540
Lys	Lys	Tyr	Tyr	Trp 545	Cys	Leu	Asp	Ser	Gly 550	Pro	Ser	Gly	Leu	Gly 555
Ile	Val	Ala	His	Met 560	Phe	Thr	Cys	Pro	Ser 565	Gly	Leu	Tyr	Phe	Asn 570
Pro	Ala	Ala	Asp	Ser 575	Cys	Asp	Phe	Ala	Arg 580	Asn	Val	Pro	Cys	Lys 585
Thr	Lys	Lys	Ser	Thr 590	Thr	Ala	Ala	Pro	Val 595	Thr	Ser	Thr	Thr	Pro 600

Ala	Thr	Thr	Thr	Val 605	Arg	Ser	Asn	Arg	Val 610	Thr	Ala	Ala	Pro	Thr 615
Ser	Arg	Pro	Val	Tyr 620	Pro	Arg	Thr	Thr	Thr 625	Thr	Thr	Ser	Thr	Thr 630
Thr	Thr	Thr	Thr	Thr 635	Thr	Pro	Ser	Thr	Val 640	Asp	Glu	Asp	Leu	Glu 645
Tyr	Glu	Glu	Asp	Thr 650	Asp	Glu	Leu	Ser	Pro 655	Ser	Lys	Ser	Thr	Asp 660
Ala	Glu	Glu	Asp	Pro 665	Gln	Val	Ile	Lys	Glu 670	Leu	Ile	Asp	Leu	Ile 675
Arg	Lys	Val	Gly	Gly 680	Val	Glu	Gln	Leu	Glu 685	Lys	His	Leu	Leu	Arg 690
Asn	Lys	Asp	Gly	Ser 695	Ile	Thr	Leu	Lys	Glu 700	Asn	Ser	Ala	Thr	Gly 705
Ala	Ala	Thr	Thr	Pro 710	Ser	Thr	Ile	Ser	Lys 715	Ser	Leu	Tyr	Asp	Arg 720
Val	Leu	Ser	Arg	Pro 725	Gly	Thr	Leu	Asn	Ser 730	Phe	Ser	Arg	Asn	Arg 735
Phe	Lys	Ile	Ser	Glu 740	Ala	Ser	Glu	Thr	Ser 745	Thr	Glu	Pro	Thr	Thr 750
Ser	Ser	Ser	Ser	Ser 755	Arg	Gly	Ser	Ser	Thr 760	Leu	Thr	Ser	Asn	Thr 765
Asn	Ser	Lys	Tyr	Ser 770	Ser	Val	Leu	Arg	Gly 775	Asn	Ser	Arg	Gln	Gly 780
Pro	Gln	Asn	Glu	Gly 785	Ile	Glu	Lys	Leu	Ala 790	Glu	Phe	Asp	Gly	Phe 795
Leu	Lys	Glu	Arg	Lys 800	Gln	Tyr	Val	Thr	Ile 805	Asn	Arg	His	Arg	Ser 810
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Glu	Glu	Asn	Leu	Ala 830	Glu	Val	Glu	Thr	Thr 835	Thr	Arg	Arg	Pro	Leu 840
Ser	Ser	Ile	Thr	Pro 845	Ser	Tyr	Thr	Ser	Leu 850	Arg	Arg	Ser	Arg	Pro 855
Thr	Thr	Val	Ala	Pro 860	Pro	Ala	Glu	Glu	Ser 865	His	Glu	Glu	Ala	Glu 870
Gln	Gln	Thr	Gln	Thr 875	Gln	Val	Lys	Ser	Tyr 880	Ala	Thr	Leu	Ser	Arg 885
Thr	Arg	Gly	Arg	Thr 890	Thr	Ser	Ser	Pro	Glu 895	Val	Thr	Glu	Ala	Ala 900
Pro	Ser	Ser	Thr	Thr	Asn	Arg	Tyr	Lys	Tyr	Phe	Glu	Arg	Thr	Arg

		905				9	910					915
Pro Th	nr Lys	Ser Ala 920	Thr	Ala	Glu		Ser 925	Glu	Asp	Pro	Thr	Glu 930
Asp Gl	lu Glu	Glu Glu 935	Tyr	Glu	Asp		Gln 940	Lys	Asp	Ile	Val	Thr 945
Val G	ln Ser	Lys Gln 950	Ser	Thr	Asn		Arg 955	Lys	Tyr	Ala	Ser	Ile 960
Gly Ar	rg Arg	Thr Thr 965	Thr	Thr	Thr		Ala 970	Thr	Pro	Glu	Thr	Thr 975
Thr Th	nr Thr	Thr Thr 980	Thr	Thr	Ala		Thr 985	Glu	Thr	Ala	Lys	Ala 990
Ser Th	nr Thr	Thr Asn 995	Asn	Asn	Asn		Asn 000	Asn	Ser	His	_	Asn .005
Ser Se	er Asn	Asn Asn 1010	Asn	Asn	Val		Leu )15	Asn	Asn	Gln		Pro .020
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Thr Se	er Thr	Thr Thr 1055	Thr	Ser	Ile		Asn 060	Asn	Leu	His		Thr .065
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Ser Gl	lu Ala	Ser Pro 1100	Thr	His	Leu		Pro 105	Ser	Pro	Asp		Glu 110
Thr Se	er Thr	Pro Thr 1115	Thr	Thr	Ser		hr 120	Thr	Thr	Glu		Pro 125
Glu Le	eu Asp	Thr Thr 1130	Thr	Thr	Thr		.ys .35	Thr	Thr	Thr		Thr 140
Thr Th	r Gly	Asn Asn 1145	Glu	Leu	Asn	_	/al .50	Asn	Asn	Val		Glu 155
Asp Se	er Glu	Val Thr 1160	Lys	Thr	Lys		51n .65	Tyr	Lys	Tyr		Thr 170
Thr As	sn Arg	Arg Arg 1175	Ile	Thr	Thr		hr .80	Thr	Thr	Ala		Lys 185
Asn Se	er Asn	Asn Asn 1190	Asn	Asn	Ala		1la .95	Ala	Asn	Asp		Ser 200
Pro Th	r Thr	Äsn Gly 1205	Leu	Ser	Ser		Asn 210	Ser	Ile	Arg		Asn 215

Pro Gly Arg Arg Gln Pro Gln Pro Glu Gln Thr Gln Thr Thr 1225 Ser Glu Pro Asn Leu Ser Ser Pro Arg Pro Phe Gly Tyr Pro Arg Arg Arg Thr Arg Pro Thr Val Ser Thr Thr Thr Thr Ile Ser 1250 Gln Thr Asp Asn Asp Asn Asn Thr Asp Asn Asp Asn Glu Thr 1265 1270 Asp Ala Val Ala Gln Val Val Lys Lys Thr Arg Leu Ser Pro Gly 1280 1285 Asp Arg Pro Lys Val Ser Ala Ser Leu Pro Thr Ala Thr Ala Ile 1295 1300 Asn Thr Arg Thr Asn Thr Ser Ser Leu His His Gln Glu Ser Gln 1310 Val Glu Val Ala Gly Asn Gly Gly Asn Asp Ser Leu Arg His Asp Val Val Ser Ser Ser Leu Ser Gln Ser Gln Ser Asn Lys Ile Asp Thr Asp Asp Leu Ser Thr Thr Gln Gln His Thr Lys Tyr Thr Trp 1355 1360 Arg Ala Val Arg Arg Pro Ala Ser Gln Arg Thr Val Val Pro Asn Ser Leu Ala Gly Asp Asp Lys Asp Ser Arg Arg Phe Ala Gly Lys 1385 1390 Gln Leu Asn Thr Glu Ser Ile Val Asp Asp Glu Leu Gln Thr Thr 1400 1405 Thr Lys Phe Arg Ser Arg Arg Leu Asn Ser Ala Glu Asp Glu Ser 1415 1420 Glu Val Ala Leu Glu Val Ala Thr Ala Thr Pro Thr His Gly Ser 1430 1435 Arg Ser Tyr Gln Ser Ile Gln Arg Ser Ala Ser Lys Ala Ser Leu 1445 1450 1455 Asp Asp Ser Gln Ile His Tyr Lys Ala Ile Ile Arg Asp Ser Glu 1460 1465 Gly Gly Ala His Leu Thr Ala Gly Arg Ser Ser Phe Val Arg 1475 1480 Asn Phe Gly Asp Ala Ala Lys Pro Thr Pro Pro His Gln Pro Ile 1495 Ser Arg Gly Gln Ile Val Glu Ser Thr Thr Glu Asp Glu Asn 1510

Val Ala Ala Glu Ile Ile Asp Asp Glu Lys Arg Gly Glu Thr Lys 1520 Ala Pro Ala Gly Ser Glu Asn Thr Asp Asp Ser Asn Thr Ala Thr 1535 Glu Gln Glu Ser Pro Glu Ile Val Thr Glu Ala Ala Gln Pro Gln 1555 Leu Glu Ile Thr Thr Leu Pro Ser Glu Thr Ser Asp Val Ser Ser 1565 1570 Ser Thr Glu Gln Ser Val Ser Ser Thr Thr Glu Glu Ser Ser Ser 1580 1585 Ser Thr Ala Asp Leu Asp Ile Val Ala Glu Glu Ala Ser Leu Gly 1595 1600 Ala Glu Thr Asp Lys Lys Ser Thr Ser Glu Asn Asp Asn Gly Glu 1610 1615 Ser Ser Thr Glu Ile Ser Ser Ser Glu Ala Pro Ile Ser Ser Thr Thr Gly Gln Ser Glu Asp Val Ser Ser Thr Thr Glu Thr Asn Ser 1645 Glu Ala Ile Glu Lys Glu Ile Ala Ser Asp Ser Asn Asp Gly Ser Ser Asp Asp Pro Ala Ser Ser Thr Glu Phe Ile Glu Ile Thr Asn 1675 Thr Thr Ser Ser Pro Val Ser Leu Gln Glu Asp Ser Ser Thr Thr Thr Glu Lys Leu Thr Arg Arg Ala Phe Asn Arg Phe Ser Ser Thr 1700 1705 Thr Pro Ala Val Val Pro Glu Asp Glu Thr Thr Ser Thr Val Asn 1715 1720 Gln Arg Arg Val Ile Val Arg Asn Arg Ile Ser Thr Thr Glu 1730 1735 1740 Ala Glu Ser Glu Ala Gln Thr Thr Glu Glu Pro Lys Arg Arg 1745 1750 Ser Phe Tyr Arg Thr Ser Thr Thr Ala Glu Pro Ser Ser Ser Thr 1760 1765 1770 Glu Ala Asp Ser Asp Ala Gln Ile Ser Thr Glu Thr Thr Arg 1775 Arg Ser Phe Phe Arg Thr Arg Thr Thr Glu Ala Ala Ser Ser Thr 1790 Thr Glu Glu Pro Ser Ser Pro Thr Glu Pro Glu Ile Glu Val Glu

Thr Thr Glu Gly Pro Thr Arg Arg Ser Phe Phe Arg Arg Ser

1820	1825	1830
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- Thr Thr Val Ala Pro Ser Ser Thr Thr Glu Glu Ile Ser Ser Ser 1835
- Ser Val Asp Asp Asp Ala Glu Ala Asn Ile Ile Thr Thr Arg Arg 1850 1855 1860
- Ser Leu Phe Thr Thr Pro Ala Pro Ser Ser Thr Glu Ala Thr Thr
  1865 1870 1875
- Thr Ala Thr Ala Glu Asp Ser Glu Val Ser Ser Ser Thr Arg Arg 1880 1885 1890
- Ser Phe Phe Arg Thr Ser Thr Thr Glu Gly Thr Thr Ser Thr
  1895 1900 1905
- Thr Glu Glu Ala Lys Asp Ile Glu His Glu Ser Glu Thr Thr Ala 1910 1915 1920
- Ala Leu Pro Lys Arg Arg Val Ile Val Arg Gly Asn Phe Arg Pro 1925 1930 1935
- Arg Lys Glu Gly Asp Leu Ser Ser Leu Leu Ala Ala Asp Ala Asn 1940 1945 1950
- Lys Arg Val Arg Asn Asn His Ser Thr Thr Ser Thr Glu Thr Pro 1955 1960 1965
- Ala Asn Ser Gln Ser Thr Thr Ser Asn Glu Glu Asp Thr Val Ala 1970 1975 1980
- Gln Pro Pro Gln Ala Glu Val Lys Ala Thr Thr Gly Arg Val Ser 1985 1990 1995
- Leu Asn Ala Val Arg Asn Arg Thr Thr Thr Lys Thr Glu Ser Leu 2000 2005 2010
- Gly Asn Gly Ile Thr Arg Thr Arg Thr Thr Tyr Val Arg Thr Leu 2015 2020 2025
- Asp Ala Gly Gln Lys Ile Val Lys Arg Ile His Thr Lys Thr Ile 2030 2035 2040
- Glu Glu Lys Pro Ala Glu Tyr Glu Tyr Ile Ile Asp Glu Val Thr 2045 2050 2055
- His Pro Pro Ala Ala Ser Thr Thr Pro Arg Thr Val Thr Arg Asn 2060 2065 2070
- Arg Gly Ser Val Arg Phe Gln Ser Asn Asp Leu Ser Ser Leu Leu 2075 2080 2085
- Ala Leu Asp Phe Ala Ser Arg Ser Thr Arg Lys Lys Gln Ala Gln 2090 2095 2100
- Thr Glu Thr Thr Val Thr Lys Thr Arg Arg Arg Leu Leu Lys Lys 2105 2110 2115
- Pro Lys Glu Thr Ile Glu His Glu Glu Val Glu Glu Tyr Glu Tyr 2120 2125 2130

Glu Ala Gly Gln Glu Ala Gly Asn Glu Val Glu Glu Ala Pro Arg Val Ser Thr Thr Ala Arg Thr Ile Ile Arg Arg Thr Arg Pro Thr Thr Ile Arg Thr Thr Thr Glu Thr Pro Gln Asn Ile Glu Ala Ser Thr Arg Arg Ala Ser Phe Ala Phe Lys Arg Pro Ser Lys Val Ser Thr Thr Glu Glu Pro Thr Thr Ser Ser Thr Glu Pro Thr Ile Ser Ala Glu Ala Thr Thr Arg Arg Val Leu Asn Phe Arg Arg Pro Val Ser Thr Thr Ser Thr Pro Ala Ser Asp Glu Ser Thr Glu Glu Ala Thr Ala Ala Pro Ile Glu Ala Thr Thr Arg Arg Val Leu Ala Phe Lys Arg Pro Val Ser Thr Thr Thr Thr Pro Ala Pro Val Asp Glu Glu Ser Thr Glu Glu Ser Thr Pro Thr Ser Ile Glu Gly Asn Thr Arg Arg Ile Leu Ala Tyr Arg Arg Pro Val Ser Thr Thr Thr Thr Pro Val Pro Val Glu Asp Glu Ser Ser Thr Asp Gln Leu Ala Ala Lys Gln Lys Phe Ile Asn Arg Leu Lys Ser Ser Thr Thr Thr Thr Ser Ile Pro Glu Thr Thr Thr Glu Glu Asp Leu Ser Asp Leu Lys Val Gln Leu Ser Asn Ala Ile Asn Arg Leu Gln Thr Glu Asn Lys Leu Glu Val Gln Thr Ile Thr Lys Gly Ser Glu Ala Ala Glu Asp Glu Gly Asp Asp Lys Leu Ser Leu Pro Ile Tyr His Arg Arg Lys Tyr Tyr Gln Tyr Val Lys Asp Ser Pro Ile Thr Tyr Ile Asp Lys Ser Pro Ala Pro Pro Asp Ile Glu Ser 

Val Thr Val Asn Ile Lys Gln Gln Ile His Asp Val Phe Asn Val

Ser Glu Asn Glu Thr Pro His Asn Ser Leu Gly Asp Asp Glu Glu 2435 2440 Thr Glu Gly His Arg Val Ala Met Ala Gln Ala Lys Glu Ile Asn 2450 2455 Ala Glu Leu Glu Glu Lys Glu Arg Gly Glu Asp Glu Ala Arg Ala 2465 2470 Leu Arg Thr Tyr Thr Arg Leu Asn Arg Thr Arg Leu Thr Leu Ser 2480 2485 Thr Arg Leu Gln Glu Lys Thr Gln Ser Glu Pro Leu Asp Thr Thr Thr Arg Arg Ser Tyr Ser Val Pro Gln Arg Phe Arg Ile Arg Ser Thr Thr Pro Ile Pro Ser Lys Ile Glu Asn Ser Glu Glu Asp Asp 2530 Glu Glu Thr Lys Asp Asn Glu Gly Pro Ser Pro Ser Thr Thr 2540 Val Thr Pro Pro Ser Ile Lys Leu Pro Thr Arg Arg Leu Phe Thr 2555 2560 Pro Arg Arg Pro Val Asn Ala Val Glu Asp Ser Asp Ser Ser Asp 2570 Ile Arg Lys Asp Asn Glu Glu Glu Leu Lys Val Glu Ser Thr Thr 2590 Lys Arg Leu Tyr Ala Gly Leu Asn Arg Leu Arg Gly Arg Gly Ser Thr Thr Thr Thr Glu Glu Ala Thr Asp Ser Thr Thr Glu Thr Ala Thr Thr Thr Ala Lys Ser Thr Arg Gln Pro Tyr Val Gly Ile 2635 Ser Arg Arg Val Thr Thr Thr Thr Thr Glu Lys Ser Ala Glu 2645 2650 Ser Ser Thr Glu Tyr Asn Gly Asn Glu Asp Glu Glu Thr Glu Ser 2665 Thr Thr Val Thr Pro Glu Gln Glu Ile Ser Asp Asp Ala Glu Glu 2675 2680 2685 Asn Lys Val Ala Ile Lys Glu Ile Asp Asp Gln Val Ser Lys Lys 2690 2695 Ala Pro Glu Glu Ala Glu Asp Thr Ser Thr Glu Glu Pro Glu Leu 2705 2710 Glu Ala Phe Ile Asp Asp Asp Asn Glu Ile Pro Leu Glu Glu Ser

Gly Pro Lys Thr Glu Thr Thr Ser Thr Thr Thr Thr Thr Ser

	2735		2740			2745
Thr Thr Thr Thr	Thr Pro 2750	Ala Ser	Thr Thr 2755	Ser Ar	g Arg	Gln Leu 2760
Val Ile Arg Arg	g Arg Phe 2765	Asn Gly	Thr Ile 2770	Thr Th	r Thr	Thr Thr 2775
Val Ala Pro Val	Ala Asp 2780	Glu Asr	Leu Glu 2785	Asn Gl	ı Ile	Asp Pro 2790
Ser Asp Thr Glu	Ser Ser 2795	Thr Pro	Lys Ala 2800	Ala Th	r Thr	Thr Ser 2805
Pro Arg Arg Glr	Leu Leu 2810	Ile Arg	Arg Arg 2815	Phe Ası	n Ala	Thr Ser 2820
Ser Gly Ser Thr	Thr Thr 2825	Thr Thr	Ala Asn 2830	Pro Sei	c Ala	Asp Asn 2835
Glu Ile Asp Glr	Gly Glu 2840	Thr Lys	Arg Thr 2845	Thr Ar	g Arg	Pro Ile 2850
Leu Ser Arg Arg	Arg Phe 2855	Asn Ala	Thr Ser 2860	Ile Th	Ala	Thr Thr 2865
Thr Gly Ser Thr	Asn Gly 2870	Asp Glu	Ile Ser 2875	Thr Arg	g Arg	Pro Tyr 2880
Ala Ala Leu Asr	a Arg Ser 2885	Arg Asn	Arg Phe 2890	Thr Thi	r Pro	Gln Thr 2895
Thr Thr Thr Asp	Gly Gly 2900	Ala Asn	Gly Asp 2905	Asp Asp	Asp	Tyr Asp 2910
Gly Glu Glu Glu	ı Glu Gln 2915	Leu Ala	Pro Pro 2920	Arg Ala	a Val	Phe Leu 2925
Gln Thr Asn Arg	His Arg 2930	Ala Leu	Lys Pro 2935	Thr Pro	Glu	Asp Glu 2940
Glu Glu Gly Ala	Ala Ala 2945	Val Pro	Gly Arg 2950	Arg Pro	Leu	Asn Phe 2955
Ala Ala Arg Arg	Thr Thr 2960	Ala Ala	Pro Leu 2965	Arg Val	. Ser	Ser Ser 2970
Thr Arg Arg Asn	Leu Val 2975	Ala Ile	Asn Arg 2980	Asn Leu	ı Tyr	His Arg 2985
Pro Glu Glu Asp	Asn Glu 2990	Glu Glu	Pro Glu 2995	Glu Glu	ı Tyr	Asp Glu 3000
Asn Glu Asp Gly	Asp Asp 3005	Asp Gln	Glu Glu 3010	Ser Val	. Asp	Pro Gln 3015
Val Thr Ser Thr	Thr Thr 3020	Arg Ser	Arg Leu 3025	Asn Glr	Leu	Leu Ala 3030
Asn Arg Gln Arg	Gln Pro 3035	Leu Arg	Thr Thr 3040	Thr Glu	Lys	Gln Thr 3045

- Glu Thr Asp Ser Asn Asp Thr Glu Thr Asp Ser Asp Asn Gly Asp 3050 3055 3060
- Glu Asn Asp Asp Glu Asp Asn Asp Ser Ser Val Glu Val Ser 3065 3070 3075
- Asn Ser Asn His Thr Leu Lys His Ser Thr Ile Phe Gly Val Gly 3080 3085 3090
- Thr Thr Asn Phe Asn Asn Leu Thr Asn Arg Ser Thr Ala Leu Asn 3095 3100 3105
- Val Ala Ser Gln Arg Ser Asn Ser Thr Val Ala Asn Tyr Ile Asn 3110 3115 3120
- Arg Phe Lys Ser Asn Ser Tyr Thr Asn Lys Asn Lys Pro Val Thr 3125 3130 3135
- Val Thr Ala Asn Ile Lys Ala Asp Ser Thr Asp Asp Lys Asp Asn 3140 3145 3150
- Tyr Ala Ser Leu Glu Asn Glu Gly Lys Glu Lys Thr Ser Gly Ala 3170 3175 3180
- Gly Leu Asn Ala Leu Gly Asn Asp Val Asn Ser Thr Arg Arg Phe 3185 3190 3195
- Gln Asn Arg Tyr Gln Leu Ser Arg Thr Arg Gly Ser Thr Thr Thr 3200 3205 3210
- Asn Thr Asn Pro Thr Thr Gln Gln Pro Gln Thr Thr Ser Thr 3215 3220 3225
- Ala Arg Arg Leu Ala Phe Gly Gly Arg Gln Arg Ala Gln Val Thr 3230 3235 3240

G. 1

- Lys Leu Thr Leu Val Asp Glu Gln Thr Glu Glu Thr Glu Thr Lys 3245 3250 3255
- Gly Asp Ser Arg Glu Glu Glu Lys Glu Glu Glu Glu Glu Glu Asp 3260 3265 3270
- Ser Asn Ala Thr Thr Thr Thr Thr Thr Thr Thr Thr Ser Arg Pro \$3275\$ \$3280\$ \$3285
- Thr Pro Lys Arg Ile Arg Val Leu Lys Phe Arg Arg Pro Leu Asn 3290 3295 3300
- Ser Asn Ser Asn Ser Thr Ile Asn Val Asp Ser Thr Thr Asn Ser 3305 3310 3315
- Ala Thr Asp Thr Asn Pro Asp Thr Thr Thr Ala Thr Pro Thr Thr 3320 3325 3330
- Ala Gly Gln Ser Thr Thr Ser Asn Ser Asn Asn Asn Asn Asn Asn and 3335 3340 3345

Thr Thr Ser Thr Thr Gly Asn Lys Arg Phe Arg Lys Ile Val Arg 3350 3355 Lys Leu Arg Pro Val Asp Ser Ser Thr Ala Ala Ser Val Asp Asn 3365 3370 3375 Ser Asp Glu Thr Thr Arg Lys Pro Phe Val Pro Ser His Thr Arg 3380 3385 Phe Ala Asp Gln Asp Asn Asp Leu Val Asn Leu Arg Gln Arg Ile 3395 Lys Glu Gln Gln Ala Arg Gly Glu Pro Gln Asp Gly Val Ile Ser Asn Arg Phe Lys Thr Leu Gly Gln Lys Asp Asp Gln Asp Val Ser 3425 3430 Glu Leu Gln Lys Leu Arg Asp Lys Val Lys Ala Glu Gln Ala Arg 3445 Gly Glu Gly Glu Gln Gly Val Ile Asn Asp Arg Leu Lys Lys Leu Leu Ala Glu Lys Gly Ser Ser Ile Ser Ser Gln Arg Glu Glu Ser 3475 Ser Thr Asp Asp Ser Ser Ser Val Ser Ser Ala Arg Pro Phe Phe Lys Arg Lys Leu Val Ala Arg Arg Pro Tyr Thr Pro Pro Ser Ala Ser Gly Gly Thr Thr Lys Ala Pro Leu Thr Phe Ser Thr Ser Arg Pro Thr Ala Lys Phe Val Arg Arg Lys Asn Gly Arg Phe Asp 3530 Pro Phe Asn Ser Ser Val Arg Asn Arg Gly Glu Gly Phe Val Arg 3550 Ser Asp Pro Arg Gly Ser Arg Leu Pro Gly Thr Asp Arg Phe Lys 3565 3570 Ser Gln Gly Asn Ser Glu Asp Asp Glu Val Glu Glu Arg His 3575 3580 Glu Gln Pro Leu Gln Asn Gln Phe Ala Thr Thr Leu Arg Arg Pro 3590 3595 3600 Phe Val Pro Lys Thr Arg Pro Val Leu Asp Lys Ser Lys Pro Glu 3605 3610 Gln Glu Asp Gly Ala Glu Glu Ser Glu Glu Glu Asp Glu Glu Glu 3620 3625 

Asp Val Lys Pro Gly Gly Glu Glu Asp Asn Ala Gln Glu Asp Asn

			3650				365	5			3660
Lys	Pro	Lys	Phe Asn 3665	Ser	Pro	Tyr	Lys Pro		Asp	Asn	Arg Ala 3675
Pro	Pro	Gly	Ser Arg 3680	Pro	Thr	Phe	Gly Th: 368		Gly	Ser	Gly Ser 3690
Pro	Pro	Thr	Ala Ser 3695	Gly	Asn	Val	Pro Ty:		Pro	Arg	Asn Arg 3705
Pro	Ser	Asn	Ser Ala 3710	Asn	Gly	Asn	Ser Th: 371		Ser	Asn	Arg Phe 3720
Gly	Thr	Thr	Lys Arg 3725	Pro	Arg	Val	Val Ası 3730	-	Pro	Pro	Gly Val 3735
Ala	Ser	Pro	Asn Leu 3740	Thr	Leu	Lys	Pro Val 3745		Ser	Asp	Tyr Glu 3750
Arg	Thr	Thr	Pro Leu 3755	Thr	Pro	Leu	Lys Pro		Pro	Phe	Ile Pro 3765
Ser	Asn	Asn	Arg Ser 3770	Tyr	Glu	Arg	Lys Tyr		Gly	Pro	Ser Thr 3780
Glu	Ala	Ala	Glu Thr 3785	Ala	Ser	Glu	Asn Ser 3790		Ile	Glu	Asp Leu 3795
Asn	Ile	Asp	Ala Leu 3800	Asn	Ala	Arg	Asn Lys		Ile	Phe	Asp Lys 3810
His	Ser	Lys	Lys His 3815	Pro	Ala	Leu	Lys Pro	_	Val	Val	Lys Val 3825
Glu	Ser	Glu	Thr Gly 3830	Leu	Glu	Val	Glu Ala 3835		Thr	Glu	Val Ala 3840
Val	Glu	Asp	Glu Thr 3845	Thr	Glu	Glu	Gln Glr 3850		Glu	Gln	Gly Phe 3855
Val	Thr	Thr	Thr Pro 3860	Ser	Thr	Pro	Pro Ser 3865		Ala	Pro	Pro Ser 3870
Thr	Gln	Ser	Asp Thr 3875	Ala	Thr	Thr	Thr Asp 3880		Pro	Pro	Glu Thr 3885
Glu	Thr	Glu	Thr Glu 3890	Thr	Glu	Thr	Glu Thr 3895		Thr	Glu	Thr Glu 3900
Asn	Val	Thr	Glu Ile 3905	Glu	Thr	Ala	Thr Asr 3910		Asn	Glu	Ala Thr 3915
Ser	Ile	Asn	Ser Gln 3920	Asp	Gln	Thr	Ile Ser 3925		Thr	Thr	Gln Ala 3930
Pro	Pro	Pro	Ala Thr 3935	Thr	Leu	Leu	His Val 3940		Thr	Leu	Leu Glu 3945
Gly	Glu	Gly	Gln Glu 3950	Glu	Glu	Pro	Thr Thr 3955	_	Lys	Pro	Thr Val 3960

64.4

Arg Leu Tyr Pro Thr Ile Gln Thr Glu Val Val Pro Lys His Lys 3965 3970 3975 Leu Ile Glu Ile Asn Arg Ile Val Glu Ile Asn Ser Lys Gln Ala 3980 3985 Lys Ala Ala Gln Arg Lys Ser Lys Ala Asn His Asp Phe Ser Thr 3995 4000 Leu Met Val Glu Ser Leu Pro His Val Glu Gln Leu Gly Glu Ile .4015 Ser Val Val Lys Tyr Val His Leu Val Asp Gly Ser Asp Ile Gln 4025 Ile Asn Asp Gly His Ser Thr Val Ala Asp Tyr Thr Pro Thr Glu Pro Thr Ser Ala Ala Glu Arg Pro Val Ser Leu Pro Val Arg Asn Ser Leu Pro Glu Thr Glu Gly Ala Asp Thr Asp Arg Ser Gly Lys Ser Leu Val Pro Glu Val Leu Thr Ala Ala Leu Glu Thr Ser Thr 4085 Ile Ser Leu Glu Gly Leu Phe Asp Ser Ala Arg Lys Gly Lys Gln Leu Ser Ser Asn Thr Ile Ile Gly Glu Thr Glu Glu Ser Thr Thr 4115 Ile Gly Ser Ser Ser Leu Ala Ser Glu Thr Gly Glu Thr Thr 4135 Thr Pro Ala Pro Thr Tyr Val Arg Pro Ile Val Pro Leu Leu Arg 4145 Pro Glu Ser Asn Glu Ser Ser Pro Leu Val Ile Ser Ile Ala Asn 4160 4165 Leu Asp Gln Val Ile Leu Ser Lys Val Gln Lys Ser Leu Ala Glu 4175 4180 Asn Ser Gln Thr Thr Val Ala Pro Glu Ala Ala Ser Asp Ser Asn 4190 4195 Ser Ala Phe Ser Val Arg Gln Pro Leu Val Val Gln Ala Pro Ile 4205 4210 Ser Asn Gly Ala Gln Glu Ile Asp Gln Asp Thr Leu Asn Thr Gln 4220 4225 4230 Asp Gln Thr Ile Asn Gly Ala Ile Ser Val Lys Thr Asn Pro Ile 4235 4240 Ile Gln Thr Thr Asn Arg Pro Asn Asp Asp Gln Val Ala Glu 4250 4255

Glu Thr Thr Ile Phe Ser Ile Glu Thr Ala Thr Glu Pro Glu Leu
4265 4270 4270

Asn Thr Gln Thr Thr Ile Pro Lys Thr Glu Ala Asn Ser Glu Thr 4280 4285 4290

Val Thr Ala Met Pro Ile Gly Ala Val Ile Met Gly Gln Phe Gly 4295 4300 4305

Leu Asn Thr Gln Ser Thr Thr Ala Val Asp Asn Asp Asn Gln Leu
4310 4315 4320

Asn Ala Gln Thr Thr Ser Thr Ile Ser Ser Gly Ala Val Ser Ser 4325 4330 4335

Val Ala Ile Gly Gly Asn Thr Gln Thr Ala Asn Ala Asp Asn Ala 4340 4345 4350

Arg Gln Glu Asn Thr Gln Ser Thr Gly Thr Ile Thr Ser Glu Ile 4355 4360 4365

Ser Ser Gly Ala Ile Ser Ser Asp Asn His Asn His Ile Gly Thr 4370 4375 4380

Gln Thr Thr Ala Thr Ile Asp Ser Ser Ser Glu Thr Thr Pro Thr 4385 4390 4395

Gln Ile Ser Thr Thr Ile Ser Ser Gly Ala Ile Ser Gly His Ile
4400 4405 4410

Asp Gly Ser Ile Asn Leu Asn Thr Gln Thr Asn Thr Thr Ile Ser 4415 4420 4425

Thr Asn Asn Thr Thr Ser Thr Thr Asp Val Gly Ser Lys Val 4430 4435 4440

Ser Glu Ala Val Ser Phe Ser Ser Glu Thr His Val Val His Arg 4445 4450 4450

Lys Lys Met Gly Arg Lys Gly Arg Gly Arg Arg Leu Arg Asn Arg
4460 4465 4470

Lys Thr Thr Thr Thr Thr Thr Thr Glu Thr Pro Thr Thr Thr 4475 4480 4485

Glu Ala Thr Phe Asp Asp Thr Thr Thr Val Val Pro Glu 4490 4495

<210> 69

<211> 782

<212> DNA

<213> Homo sapien

<400> 69

aggggcctta gcgtgccgca tcgccgagat ccagcgcca gagagacacc 50
agagaaccca ccatggcccc ctttgagccc ctggcttctg gcatcctgtt 100
gttgctgtgg ctgatagccc ccagcagggc ctgcacctgt gtcccacccc 150
acccacagac ggccttctgc aattccgacc tcgtcatcag ggccaagttc 200

qtqqqqacac caqaaqtcaa ccaqaccacc ttataccaqc qttatqaqat 250 caagatgacc aagatgtata aagggttcca agccttaggg gatgccgctg 300 acateeggtt egtetacace ecegecatgg agagtgtetg eggatactte 350 cacaggtccc acaaccgcag cgaggagttt ctcattgctg gaaaactgca 400 ggatggactc ttgcacatca ctacctgcag tttcgtggct ccctggaaca 450 geetgagett ageteagege eggggettea ecaagaceta eactgttgge 500 tgtgaggaat gcacagtgtt tccctgttta tccatcccct gcaaactgca 550 gagtggcact cattgcttgt ggacggacca gctcctccaa ggctctgaaa 600 agggetteca gtecegteae ettgeetgee tgeeteggga geeagggetg 650 tgcacctggc agtccctgcg gtcccagata gcctgaatcc tgcccggagt 700 ggaactgaag cctgcacagt gtccaccctg ttcccactcc catctttctt 750 ccggacaatg aaataaagag ttaccaccca gc 782

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<210> 70
<211> 207
<212> PRT
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<213> Homo sapien

<400> 70

Met Ala Pro Phe Glu Pro Leu Ala Ser Gly Ile Leu Leu Leu Trp Leu Ile Ala Pro Ser Arg Ala Cys Thr Cys Val Pro Pro His Pro Gln Thr Ala Phe Cys Asn Ser Asp Leu Val Ile Arg Ala Lys Phe Val Gly Thr Pro Glu Val Asn Gln Thr Thr Leu Tyr Gln Arg Tyr Glu Ile Lys Met Thr Lys Met Tyr Lys Gly Phe Gln Ala Leu Gly Asp Ala Ala Asp Ile Arg Phe Val Tyr Thr Pro Ala Met Glu Ser Val Cys Gly Tyr Phe His Arg Ser His Asn Arg Ser Glu Glu 95 100 Phe Leu Ile Ala Gly Lys Leu Gln Asp Gly Leu Leu His Ile Thr 110 Thr Cys Ser Phe Val Ala Pro Trp Asn Ser Leu Ser Leu Ala Gln 125 Arg Arg Gly Phe Thr Lys Thr Tyr Thr Val Gly Cys Glu Glu Cys

140

Thr Val Phe Pro Cys Leu Ser Ile Pro Cys Lys Leu Gln Ser Gly 155 160 Thr His Cys Leu Trp Thr Asp Gln Leu Leu Gln Gly Ser Glu Lys 170 180 Gly Phe Gln Ser Arg His Leu Ala Cys Leu Pro Arg Glu Pro Gly Leu Cys Thr Trp Gln Ser Leu Arg Ser Gln Ile Ala <210> 71 <211> 481 <212> DNA <213> Homo sapien <400> 71 ccactgcacg gtagggggtc ctgtaggagg ctggtggcag ggttggattg 50 tgggccctag gcttctgggc gggatgatga cattgagatt ctggccctg 100 tatccacagg tgatggagac ctgccagatg tccaggagcc cccgagagcg 150 gctgttgctg cttttgctgc tgctactgct tgtgccctgg ggcactggcc 200 ctgcctcagg tgttgccctg ccctcgctg gtgtgttcag cctccgcgcc 250 ccgggtcgtg cctgggcggg cttgggtagc cccctgtctc ggcgcagcct 300 ggegetaget gaegaegegg cettteggga gegegegeg etgetggeeg 350 ccctggagcg ccgccgctgg ctggactctt acatgcagaa gctgttgcta 400 ctggacgege cetgageeta ataaagagee tgtegeactg egactgegee 450 tctttgctgc gccactctct tgtgggtgtg t 481

<210> 72

<211> 100

<212> PRT

<213> Homo sapien

<400> 72

Met Glu Thr Cys Gln Met Ser Arg Ser Pro Arg Glu Arg Leu Leu  $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$ 

Leu Leu Leu Leu Leu Leu Val Pro Trp Gly Thr Gly Pro 20 25 30

Ala Ser Gly Val Ala Leu Pro Leu Ala Gly Val Phe Ser Leu Arg
35 40 45

Ala Pro Gly Arg Ala Trp Ala Gly Leu Gly Ser Pro Leu Ser Arg
50 55 60

Arg Ser Leu Ala Leu Ala Asp Asp Ala Ala Phe Arg Glu Arg Ala 65 70 75

Arg Leu Leu Ala Ala Leu Glu Arg Arg Arg Trp Leu Asp Ser Tyr 80 85 90

### Met Gln Lys Leu Leu Leu Asp Ala Pro 95

<210> 73 <211> 2974

<212> DNA

<213> Homo sapien

<400> 73

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agaacgaatt aagtgttgac cacagcgacc cagtcatcct gaatgtcctc 1350 tatggcccag acgaccccac catttccccc tcatacacct attaccgtcc 1400 aggggtgaac ctcagcctct cctgccatgc agcctctaac ccacctgcac 1450 agtattettg getgattgat gggaacatee agcaacacae acaagagete 1500 tttatctcca acatcactga gaagaacagc ggactctata cctgccaggc 1550 caataactca gccagtggcc acagcaggac tacagtcaag acaatcacag 1600 tetetgegga getgeecaag ceetecatet ceageaacaa etecaaacee 1650 gtggaggaca aggatgctgt ggccttcacc tgtgaacctg aggctcagaa 1700 cacaacctac ctgtggtggg taaatggtca gagcctccca gtcagtccca 1750 ggctgcagct gtccaatggc aacaggaccc tcactctatt caatgtcaca 1800 agaaatgacg caagagccta tgtatgtgga atccagaact cagtgagtgc 1850 aaaccgcagt gacccagtca ccctggatgt cctctatggg ccggacaccc 1900 ccatcatttc cccccagac tcgtcttacc tttcgggagc gaacctcaac 1950 ctctcctgcc actcggcctc taacccatcc ccgcagtatt cttggcgtat 2000 caatgggata ccgcagcaac acacacaagt tctctttatc gccaaaatca 2050 cgccaaataa taacgggacc tatgcctgtt ttgtctctaa cttggctact 2100 ggccgcaata attccatagt caagagcatc acagtctctg catctggaac 2150 ttctcctggt ctctcagctg gggccactgt cggcatcatg attggagtgc 2200 tggttggggt tgctctgata tagcagccct ggtgtagttt cttcatttca 2250 ggaagactga cagttgtttt gcttcttcct taaagcattt gcaacagcta 2300 cagtctaaaa ttgcttcttt accaaggata tttacagaaa agactctgac 2350 cagagatega gaccatecta gecaacateg tgaaaceeca tetetaetaa 2400 aaatacaaaa atgagctggg cttggtggcg cgcacctgta gtcccagtta 2450 ctcgggaggc tgaggcagga gaatcgcttg aacccgggag gtggagattg 2500 cagtgagccc agatcgcacc actgcactcc agtctggcaa cagagcaaga 2550 ctccatctca aaaagaaaag aaaagaagac tctgacctgt actcttgaat 2600 acaagtttct gataccactg cactgtctga gaatttccaa aactttaatg 2650 aactaactga cagcttcatg aaactgtcca ccaagatcaa gcagagaaaa 2700 taattaattt catgggacta aatgaactaa tgaggattgc tgattcttta 2750 aatgtettgt tteecagatt teaggaaact ttttttettt taagetatee 2800

actcttacag caatttgata aaatatactt ttgtgaacaa aaattgagac 2850 atttacattt tctccctatg tggtcgctcc agacttggga aactattcat 2900 gaatatttat attgtatggt aatatagtta ttgcacaagt tcaataaaaa 2950 tctgctcttt gtataacaga aaaa 2974

<210> 74

<211> 702

<212> PRT

<213> Homo sapien

<400> 74

Met Glu Ser Pro Ser Ala Pro Pro His Arg Trp Cys Ile Pro Trp

1 5 10 15

Gln Arg Leu Leu Thr Ala Ser Leu Leu Thr Phe Trp Asn Pro
20 25 30

Pro Thr Thr Ala Lys Leu Thr Ile Glu Ser Thr Pro Phe Asn Val 35 40 45

Ala Glu Gly Lys Glu Val Leu Leu Val His Asn Leu Pro Gln 50 55 60

His Leu Phe Gly Tyr Ser Trp Tyr Lys Gly Glu Arg Val Asp Gly
65 70 75

Asn Arg Gln Ile Ile Gly Tyr Val Ile Gly Thr Gln Gln Ala Thr 80 85 90

Pro Gly Pro Ala Tyr Ser Gly Arg Glu Ile Ile Tyr Pro Asn Ala 95 100 105

Ser Leu Leu Ile Gln Asn Ile Ile Gln Asn Asp Thr Gly Phe Tyr 110 115 120

Thr Leu His Val Ile Lys Ser Asp Leu Val Asn Glu Glu Ala Thr 125 130 135

Gly Gln Phe Arg Val Tyr Pro Glu Leu Pro Lys Pro Ser Ile Ser 140 145 150

Ser Asn Asn Ser Lys Pro Val Glu Asp Lys Asp Ala Val Ala Phe 155 160 160

Thr Cys Glu Pro Glu Thr Gln Asp Ala Thr Tyr Leu Trp Trp Val 170 175 180

Asn Asn Gln Ser Leu Pro Val Ser Pro Arg Leu Gln Leu Ser Asn 185 190 195

Gly Asn Arg Thr Leu Thr Leu Phe Asn Val Thr Arg Asn Asp Thr 200 205 210

Ala Ser Tyr Lys Cys Glu Thr Gln Asn Pro Val Ser Ala Arg Arg 215 220 225

Ser Asp Ser Val Ile Leu Asn Val Leu Tyr Gly Pro Asp Ala Pro 230 235 240

Thr	Ile	Ser	Pro	Leu 245	Asn	Thr	Ser	Tyr	Arg 250	Ser	Gly	Glu	Asn	Leu 255
Asn	Leu	Ser	Cys	His 260	Ala	Ala	Ser	Asn	Pro 265	Pro	Ala	Gln	Tyr	Ser 270
Trp	Phe	Val	Asn	Gly 275	Thr	Phe	Gln	Gln	Ser 280	Thr	Gln	Glu	Leu	Phe 285
Ile	Pro	Asn	Ile	Thr 290	Val	Asn	Asn	Ser	Gly 295	Ser	Tyr	Thr	Cys	Gln 300
Ala	His	Asn	Ser	Asp 305	Thr	Gly	Leu	Asn	Arg 310	Thr	Thr	Val	Thr	Thr 315
Ile	Thr	Val	Tyr	Ala 320	Glu	Pro	Pro	Lys	Pro 325	Phe	Ile	Thr	Ser	Asn 330
Asn	Ser	Asn	Pro	Val 335	Glu	Asp	Glu	Asp	Ala 340	Val	Ala	Leu	Thr	Cys 345
Glu	Pro	Glu	Ile	Gln 350	Asn	Thr	Thr	Tyr	Leu 355	Trp	Trp	Val	Asn	Asn 360
Gln	Ser	Leu	Pro	Val 365	Ser	Pro	Arg	Leu	Gln 370	Leu	Ser	Asn	Asp	Asn 375
Arg	Thr	Leu	Thr	Leu 380	Leu	Ser	Val	Thr	Arg 385	Asņ	Asp	Val	Gly	Pro 390
Tyr	Glu	Cys	Gly	Ile 395	Gln	Asn	Glu	Leu	Ser 400	Val	Asp	His	Ser	Asp 405
Pro	Val	Ile	Leu	Asn 410	Val	Leu	Tyr	Gly	Pro 415	Asp	Asp	Pro	Thr	Ile 420
									1 0					
Ser	Pro	Ser	Tyr	Thr 425	Tyr	Tyr	Arg	Pro		Val	Asn	Leu	Ser	Leu 435
				425					Gly 430				Ser Trp	435
Ser	Cys	His	Ala	425 Ala 440	Ser	Asn	Pro	Pro	Gly 430 Ala 445	Gln	Tyr	Ser		435 Leu 450
Ser Ile	Cys Asp	His Gly	Ala Asn	Ala 440 Ile 455	Ser Gln	Asn Gln	Pro His	Pro Thr	Gly 430 Ala 445 Gln 460	Gln Glu	Tyr Leu	Ser Phe	Trp	435 Leu 450 Ser 465
Ser Ile Asn	Cys Asp Ile	His Gly Thr	Ala Asn Glu	Ala 440 Ile 455 Lys 470	Ser Gln Asn	Asn Gln Ser	Pro His Gly	Pro Thr Leu	Gly 430 Ala 445 Gln 460 Tyr 475	Gln Glu Thr	Tyr Leu Cys	Ser Phe Gln	Trp	435 Leu 450 Ser 465 Asn 480
Ser Ile Asn	Cys Asp Ile Ser	His Gly Thr	Ala Asn Glu Ser	Ala 440 Ile 455 Lys 470 Gly 485	Ser Gln Asn His	Asn Gln Ser Ser	Pro His Gly Arg	Pro Thr Leu Thr	Gly 430 Ala 445 Gln 460 Tyr 475 Thr 490	Gln Glu Thr Val	Tyr Leu Cys Lys	Ser Phe Gln Thr	Trp Ile Ala	435 Leu 450 Ser 465 Asn 480 Thr 495
Ser Ile Asn Asn	Cys Asp Ile Ser	His Gly Thr Ala Ala	Ala Asn Glu Ser	Ala 440 Ile 455 Lys 470 Gly 485 Leu 500	Ser Gln Asn His	Asn Gln Ser Ser	Pro His Gly Arg	Pro Thr Leu Thr	Gly 430 Ala 445 Gln 460 Tyr 475 Thr 490 Ile 505	Gln Glu Thr Val Ser	Tyr Leu Cys Lys Ser	Ser Phe Gln Thr	Trp Ile Ala Ile	435 Leu 450 Ser 465 Asn 480 Thr 495 Ser 510

Leu Pro Val Ser Pro Arg Leu Gln Leu Ser Asn Gly Asn Arg Thr 545 550 555 Leu Thr Leu Phe Asn Val Thr Arg Asn Asp Ala Arg Ala Tyr Val 560 565 570 Cys Gly Ile Gln Asn Ser Val Ser Ala Asn Arg Ser Asp Pro Val Thr Leu Asp Val Leu Tyr Gly Pro Asp Thr Pro Ile Ile Ser Pro 595 600 Pro Asp Ser Ser Tyr Leu Ser Gly Ala Asn Leu Asn Leu Ser Cys 605 610 615 His Ser Ala Ser Asn Pro Ser Pro Gln Tyr Ser Trp Arg Ile Asn 620 625 630 Gly Ile Pro Gln Gln His Thr Gln Val Leu Phe Ile Ala Lys Ile 635 640 Thr Pro Asn Asn Gly Thr Tyr Ala Cys Phe Val Ser Asn Leu Ala Thr Gly Arg Asn Asn Ser Ile Val Lys Ser Ile Thr Val Ser 675 Ala Ser Gly Thr Ser Pro Gly Leu Ser Ala Gly Ala Thr Val Gly 685 690 Ile Met Ile Gly Val Leu Val Gly Val Ala Leu Ile 695

<210> 75

<211> 2249

<212> DNA

<213> Homo sapien

<400> 75

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<210> 76

<211> 344

<212> PRT

<213> Homo sapien

<400> 76

Met Gly Pro Pro Ser Ala Pro Pro Cys Arg Leu His Val Pro Trp
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Lys Glu Val Leu Leu Thr Ala Ser Leu Leu Thr Phe Trp Asn Pro
20 25 30

Pro Thr Thr Ala Lys Leu Thr Ile Glu Ser Thr Pro Phe Asn Val\$35\$ 40 45

Ala Glu Gly Lys Glu Val Leu Leu Leu Ala His Asn Leu Pro Gln 50 55 60

Asn Arg Ile Gly Tyr Ser Trp Tyr Lys Gly Glu Arg Val Asp Gly
65 70 75

Asn Ser Leu Ile Val Gly Tyr Val Ile Gly Thr Gln Gln Ala Thr 80 85 90

Pro Gly Pro Ala Tyr Ser Gly Arg Glu Thr Ile Tyr Pro Asn Ala 95 100 105

Ser Leu Leu Ile Gln Asn Val Thr Gln Asn Asp Thr Gly Phe Tyr 110 115 120

Thr Leu Gln Val Ile Lys Ser Asp Leu Val Asn Glu Glu Ala Thr 125 130 130

Gly Gln Phe His Val Tyr Pro Glu Leu Pro Lys Pro Ser Ile Ser 140 145 150

Ser Asn Asn Ser Asn Pro Val Glu Asp Lys Asp Ala Val Ala Phe 155 160 165

Thr Cys Glu Pro Glu Val Gln Asn Thr Thr Tyr Leu Trp Trp Val 170 175 180

Asn Gly Gln Ser Leu Pro Val Ser Pro Arg Leu Gln Leu Ser Asn 185 190 195

Gly Asn Met Thr Leu Thr Leu Leu Ser Val Lys Arg Asn Asp Ala 200 205 210

Gly Ser Tyr Glu Cys Glu Ile Gln Asn Pro Ala Ser Ala Asn Arg 215 220 225

Ser Asp Pro Val Thr Leu Asn Val Leu Tyr Gly Pro Asp Val Pro 230 235 240

Thr Ile Ser Pro Ser Lys Ala Asn Tyr Arg Pro Gly Glu Asn Leu 245 250 Asn Leu Ser Cys His Ala Ala Ser Asn Pro Pro Ala Gln Tyr Ser 260 265 270 Trp Phe Ile Asn Gly Thr Phe Gln Gln Ser Thr Gln Glu Leu Phe 275 Ile Pro Asn Ile Thr Val Asn Asn Ser Gly Ser Tyr Met Cys Gln 295 300 Ala His Asn Ser Ala Thr Gly Leu Asn Arg Thr Thr Val Thr Met 310 Ile Thr Val Ser Gly Ser Ala Pro Val Leu Ser Ala Val Ala Thr 320 325 330 Val Gly Ile Thr Ile Gly Val Leu Ala Arg Val Ala Leu Ile 335

<210> 77

<211> 1386

<212> DNA

<213> Homo sapien

<400> 77

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<210> 78

<211> 317

<212> PRT

<213> Homo sapien

<400> 78

Met Val Arg Ala Arg His Gln Pro Gly Gly Leu Cys Leu Leu 1 5 10 15

Leu Leu Cys Gln Phe Met Glu Asp Arg Ser Ala Gln Ala Gly
20 25 30

Asn Cys Trp Leu Arg Gln Ala Lys Asn Gly Arg Cys Gln Val Leu 35 40 45

Tyr Lys Thr Glu Leu Ser Lys Glu Glu Cys Cys Ser Thr Gly Arg
50 55 60

Leu Ser Thr Ser Trp Thr Glu Glu Asp Val Asn Asp Asn Thr Leu 65 70 75

Phe Lys Trp Met Ile Phe Asn Gly Gly Ala Pro Asn Cys Ile Pro 80 85 90

Cys Lys Glu Thr Cys Glu Asn Val Asp Cys Gly Pro Gly Lys Lys 95 100 105

Cys Arg Met Asn Lys Lys Asn Lys Pro Arg Cys Val Cys Ala Pro 110 115 120

Asp Cys Ser Asn Ile Thr Trp Lys Gly Pro Val Cys Gly Leu Asp 125 130 135

Gly Lys Thr Tyr Arg Asn Glu Cys Ala Leu Leu Lys Ala Arg Cys 140 145

Lys Glu Gln Pro Glu Leu Glu Val Gln Tyr Gln Gly Arg Cys Lys 155 160 165

Lys Thr Cys Arq Asp Val Phe Cys Pro Gly Ser Ser Thr Cys Val 170 175 180 Val Asp Gln Thr Asn Asn Ala Tyr Cys Val Thr Cys Asn Arg Ile 185 190 195 Cys Pro Glu Pro Ala Ser Ser Glu Gln Tyr Leu Cys Gly Asn Asp 210 Gly Val Thr Tyr Ser Ser Ala Cys His Leu Arg Lys Ala Thr Cys 215 220 225 Leu Leu Gly Arg Ser Ile Gly Leu Ala Tyr Glu Gly Lys Cys Ile Lys Ala Lys Ser Cys Glu Asp Ile Gln Cys Thr Gly Gly Lys Lys 250 255 Cys Leu Trp Asp Phe Lys Val Gly Arg Gly Arg Cys Ser Leu Cys 265 Asp Glu Leu Cys Pro Asp Ser Lys Ser Asp Glu Pro Val Cys Ala Ser Asp Asn Ala Thr Tyr Ala Ser Glu Cys Ala Met Lys Glu Ala Ala Cys Ser Ser Gly Val Leu Leu Glu Val Lys Hi's Ser Gly Ser 310 315

Cys Asn

<210> 79

<211> 3445

<212> DNA

<213> Homo sapien

<400> 79

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gatgaggtgc agaagatgag gatggctgtc attgggggtg cgatatttct 600 tcttgcaggt ctggctattt tagttgccac agcatggtat ggcaatagaa 650 tegtteaaga attetatgae eetatgaeee eagteaatge eaggtaegaa 700 tttggtcagg ctctcttcac tggctgggct gctgcttctc tctgccttct 750 gggaggtgcc ctactttgct gttcctgtcc ccgaaaaaca acctcttacc 800 caacaccaag gccctatcca aaacctgcac cttccagcgg gaaagactac 850 gtgtgacaca gaggcaaaag gagaaaatca tgttgaaaca aaccgaaaat 900 ggacattgag atactatcat taacattagg accttagaat tttgggtatt 950 gtaatctgaa gtatggtatt acaaaacaaa caaacaaaca aaaaacccat 1000 gtgttaaaat actcagtgct aaacatggct taatcttatt ttatcttctt 1050 tcctcaatat aggagggaag atttttccat ttgtattact gcttcccatt 1100 gagtaatcat actcaattgg gggaaggggt gctccttaaa tatatataga 1150 tatgtatata tacatgtttt tctattaaaa atagacagta aaatactatt 1200 ctcattatgt tgatactagc atacttaaaa tatctctaaa ataggtaaat 1250 gtatttaatt ccatattgat gaagatgttt attggtatat tttcttttc 1300 gtctatatat acatatgtaa cagtcaaata tcatttactc ttcttcatta 1350 gctttgggtg cctttgccac aagacctagc ctaatttacc aaggatgaat 1400 tettteaatt etteatgegt gecettttea tataettatt ttattttta 1450 ccataatctt atagcacttg catcgttatt aagcccttat ttgttttgtg 1500 tttcattggt ctctatctcc tgaatctaac acatttcata gcctacattt 1550 tagtttctaa agccaagaag aatttattac aaatcagaac tttggaggca 1600 aatctttctg catgaccaaa gtgataaatt cctgttgacc ttcccacaca 1650 atccctgtac tctgacccat agcactcttg tttgctttga aaatatttgt 1700 ccaattgagt agctgcatgc tgttccccca ggtgttgtaa cacaacttta 1750 ttgattgaat ttttaagcta cttattcata gttttatatc cccctaaact 1800 acctttttgt tccccattcc ttaattgtat tgttttccca agtgtaatta 1850 tcatgcgttt tatatcttcc taataaggtg tggtctgttt gtctgaacaa 1900 agtgctagac tttctggagt gataatctgg tgacaaatat tctctctgta 1950 gctgtaagca agtcacttaa tctttctacc tcttttttct atctgccaaa 2000 ttgagataat gatacttaac cagttagaag aggtagtgtg aatattaatt 2050 agtttatatt actctcattc tttgaacatg aactatgcct atgtagtgtc 2100

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<sup>&</sup>lt;210> 80

<sup>&</sup>lt;211> 211

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 80

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Leu Gly Trp Ile Gly Ala Ile Val Ser Thr Ala Leu Pro Gln Trp
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                                     25
Arg Ile Tyr Ser Tyr Ala Gly Asp Asn Ile Val Thr Ala Gln Ala
Met Tyr Glu Gly Leu Trp Met Ser Cys Val Ser Gln Ser Thr Gly
Gln Ile Gln Cys Lys Val Phe Asp Ser Leu Leu Asn Leu Ser Ser
Thr Leu Gln Ala Thr Arg Ala Leu Met Val Val Gly Ile Leu Leu
                 80
                                     85
Gly Val Ile Ala Ile Phe Val Ala Thr Val Gly Met Lys Cys Met
Lys Cys Leu Glu Asp Asp Glu Val Gln Lys Met Arg Met Ala Val
                                    115
Ile Gly Gly Ala Ile Phe Leu Leu Ala Gly Leu Ala Ile Leu Val
Ala Thr Ala Trp Tyr Gly Asn Arg Ile Val Gln Glu Phe Tyr Asp
                                    145
Pro Met Thr Pro Val Asn Ala Arg Tyr Glu Phe Gly Gln Ala Leu
Phe Thr Gly Trp Ala Ala Ser Leu Cys Leu Leu Gly Gly Ala
                                    175
Leu Leu Cys Cys Ser Cys Pro Arg Lys Thr Thr Ser Tyr Pro Thr
                                                        195
Pro Arg Pro Tyr Pro Lys Pro Ala Pro Ser Ser Gly Lys Asp Tyr
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Val

<210> 81

<211> 1233

<212> DNA

<213> Homo sapien

<400> 81

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qqcaccaaca tecteacqqc eqtqteetac etqaaaqqqc tetqqatqqa 300 gtgtgtgtgg cacagcacag gcatctacca gtgccagatc taccgatccc 350 tgctggcgct gccccaagac ctccaggctg cccgcgccct catggtcatc 400 tectgeetge tetegggeat ageetgegee tgegeegtea tegggatgaa 450 gtgcacgcgc tgcgccaagg gcacacccgc caagaccacc tttgccatcc 500 teggeggeac cetetteate etggeeggee teetgtgeat ggtggeegte 550 tectggacca ccaacgacgt ggtgcagaac ttetacaacc cgctgctgcc 600 cagcggcatg aagtttgaga ttggccaggc cctgtacctg ggcttcatct 650 cctcgtccct ctcgctcatt ggtggcaccc tgctttgcct gtcctqccag 700 gacgaggcac cctacaggcc ctaccaggcc ccgcccaggg ccaccacgac 750 cactgcaaac accgcacctg cctaccagcc accagctgcc tacaaagaca 800 ategggeece cteagtgace teggeeacge acagegggta caggetgaac 850 gactacgtgt gagtccccac agcctgcttc tcccctgggc tgctgtgggc 900 tgggtccccg gcgggactgt caatggaggc aggggttcca gcacaaagtt 950 tacttctggg caatttttgt atccaaggaa ataatgtgaa tgcgaggaaa 1000 tgtctttaga gcacagggac agagggggaa ataagaggag gagaaagctc 1050 tctataccaa agactgaaaa aaaaaatcct gtctgttttt gtatttatta 1100 tatatattta tgtgggtgat ttgataacaa gtttaatata aagtgacttg 1150 ggagtttggt cagtggggtt ggtttgtgat ccaggaataa accttgcgga 1200 tgtggctgtt tatgaaaaaa aaaaaaaaaa aaa 1233

<210> 82

<211> 239

<212> PRT

<213> Homo sapien

## <400> 82

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Leu Gly Met Val Gly Thr Leu Ile Thr Thr Ile Leu Pro His Trp 20 25 30

Arg Arg Thr Ala His Val Gly Thr Asn Ile Leu Thr Ala Val Ser 35 40 45

Tyr Leu Lys Gly Leu Trp Met Glu Cys Val Trp His Ser Thr Gly 50 55

Ile Tyr Gln Cys Gln Ile Tyr Arg Ser Leu Leu Ala Leu Pro Gln 65 70 75

Asp Leu Gln Ala Ala Arg Ala Leu Met Val Ile Ser Cys Leu Leu 85 Ser Gly Ile Ala Cys Ala Cys Ala Val Ile Gly Met Lys Cys Thr 105 Arg Cys Ala Lys Gly Thr Pro Ala Lys Thr Thr Phe Ala Ile Leu 110 115 Gly Gly Thr Leu Phe Ile Leu Ala Gly Leu Leu Cys Met Val Ala 130 125 135 Val Ser Trp Thr Thr Asn Asp Val Val Gln Asn Phe Tyr Asn Pro Leu Leu Pro Ser Gly Met Lys Phe Glu Ile Gly Gln Ala Leu Tyr 165 Leu Gly Phe Ile Ser Ser Ser Leu Ser Leu Ile Gly Gly Thr Leu Leu Cys Leu Ser Cys Gln Asp Glu Ala Pro Tyr Arg Pro Tyr Gln Ala Pro Pro Arg Ala Thr Thr Thr Ala Asn Thr Ala Pro Ala 205 210 Tyr Gln Pro Pro Ala Ala Tyr Lys Asp Asn Arg Ala Pro Ser Val 220 225 Thr Ser Ala Thr His Ser Gly Tyr Arg Leu Asn Asp Tyr Val 230 235

<210> 83

<211> 4716

<212> DNA

<213> Homo sapien

# <400> 83

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<211> 1358

<212> PRT

<213> Homo sapien

<400> 84

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Gln	Pro	Val	Val	Phe 65	Asn	His	Val	Tyr	Asn 70	Ile	Asn	Val	Pro	Leu 75
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Val	Ser	Ala	Glu	Asp 95	Glu	Thr	Leu	Ala	Glu 100	Tyr	Met	Gly	Gln	Thr 105
Ser	Asp	His	Glu	Ser 110	Gln	Val	Thr	Phe	Thr 115	His	Arg	Ile	Asn	Phe 120
Pro	Lys	Lys	Ala	Cys 125	Pro	Cys	Ala	Ser	Ser 130	Ala	Gln	Val	Leu	Gln 135
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Leu	Arg	Asp	Gln	Cys 155	Asn	Ala	Asn	Cys	Cys 160	Gln	Glu	Ser	Ala	Ala 165
Thr	Gly	Gln	Leu	Asp 170	Tyr	Ile	Pro	His	Cys 175	Ser	Gly	His	Gly	Asn 180
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Gly	Lys	Asn	Cys	Ser 200	Glu	Pro	Tyr	Cys	Pro 205	Leu	Gly	Cys	Ser	Ser 210
Arg	Gly	Val	Cys	Val 215	Asp	Gly	Gln	Cys	Ile 220	Cys	Asp	Ser	Glu	Tyr 225
Ser	Gly	Asp	Asp	Cys 230	Ser	Glu	Leu	Arg	Cys 235	Pro	Thr	Asp.	Cys	Ser 240
Ser	Arg	Gly	Leu	Cys 245	Val	Asp	Gly	Glu	Cys 250	Val	Cys	Glu	Glu	Pro 255
Tyr	Thr	Gly	Glu	Asp 260	Cys	Arg	Glu	Leu	Arg 265	Cys	Pro	Gly	Asp	Cys 270
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Gly	Tyr	Val	Gly	Glu 290	Asp	Cys	Gly	Gln	Arg 295	Gln	Cys	Leu	Asn	Ala 300
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Pro Il	e Th	r Ala	Lys 410	Val	Ala	Thr	His	Leu 415	Ser	Thr	Pro	Gln	Gly 420
Leu Gl	n Ph	e Lys	Thr 425	Ile	Thr	Glu	Thr	Thr 430	Val	Glu	Val	Gln	Trp 435
Glu Pr	o Ph	e Ser	Phe 440	Ser	Phe	Asp	Gly	Trp 445	Glu	Ile	Ser	Phe	Ile 450
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Pro	Pro	Val	Ala	Ser 890	Phe	Asp	Tyr	Tyr	Arg 895	Val	Ser	Tyr	Arg	Pro 900
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Ile	Ser	Leu	Asn	Ser 935	Val	Arg	Gly	Arg	Glu 940	Glu	Ser	Glu	Arg	Ile 945
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Asn Gly Thr Ala Gly Asp Ser Leu Ser Tyr His Gln Gly Arg Pro

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Ala Met Ser Tyr Lys Gly Ala Trp Trp Tyr Lys Asn Cys His Arg 1295 1300 1305

Thr Asn Leu Asn Gly Lys Tyr Gly Glu Ser Arg His Ser Gln Gly
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Ile Asn Trp Tyr His Trp Lys Gly His Glu Phe Ser Ile Pro Phe 1325 1330 1335

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<211> 3205

<212> DNA

<213> Homo sapien

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<212> PRT

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Phe Arg Glu Ala Glu Val Thr Leu Glu Ala Gly Gly Ala Glu Gln 35 40 45

Glu Pro Gly Gln Ala Leu Gly Lys Val Phe Met Gly Cys Pro Gly
50 55 60

Gln Glu Pro Ala Leu Phe Ser Thr Asp Asn Asp Asp Phe Thr Val
65 70 75

Arg Asn Gly Glu Thr Val Gln Glu Arg Arg Ser Leu Lys Glu Arg 80 85 90

Asn Pro Leu Lys Ile Phe Pro Ser Lys Arg Ile Leu Arg Arg His 95 100 105

Lys	Arg	Asp	Trp	Val 110	Val	Ala	Pro	Ile	Ser 115	Val	Pro	Glu	Asn	Gly 120
Lys	Gly	Pro	Phe	Pro 125	Gln	Arg	Leu	Asn	Gln 130	Leu	Lys	Ser	Asn	Lys 135
Asp	Arg	Asp	Thr	Lys 140	Ile	Phe	Tyr	Ser	Ile 145	Thr	Gly	Pro	Gly	Ala 150
Asp	Ser	Pro	Pro	Glu 155	Gly	Val	Phe	Ala	Val 160	Glu	Lys	Glu	Thr	Gly 165
Trp	Leu	Leu	Leu	Asn 170	Lys	Pro	Leu	Asp	Arg 175	Glu	Glu	Ile	Ala	Lys 180
Tyr	Glu	Leu	Phe	Gly 185	His	Ala	Val	Ser	Glu 190	Asn	Gly	Ala	Ser	Val 195
Glu	Asp	Pro	Met	Asn 200	Ile	Ser	Ile	Ile	Val 205	Thr	Asp	Gln	Asn	Asp 210
His	Lys	Pro	Lys	Phe 215	Thr	Gln	Asp	Thr	Phe 220	Arg	Gly	Ser	Val	Leu 225
Glu	Gly	Val	Leu	Pro 230	Gly	Thr	Ser	Val	Met 235	Gln	Val	Thr	Ala	Thr 240
Asp	Glu	Asp	Asp	Ala 245	Ile	Tyr	Thr	Tyr	Asn 250	Gly	Val	Val	Ala	Tyr 255
Ser	Ile	His	Ser	Gln 260	Glu	Pro	Lys	Asp	Pro 265	His	Asp	Leu	Met	Phe 270
Thr	Ile	His	Arg	Ser 275	Thr	Gly	Thr	Ile	Ser 280	Val	Ile	Ser	Ser	Gly 285
Leu	Asp	Arg	Glu	Lys 290	Val	Pro	Glu	Tyr	Thr 295	Leu	Thr	Ile	Gln	Ala 300
Thr	Asp	Met	Asp	Gly 305	Asp	Gly	Ser	Thr	Thr 310	Thr	Ala	Val	Ala	Val 315
Val	Glu	Ile	Leu	Asp 320	Ala	Asn	Asp	Asn	Ala 325	Pro	Met	Phe	Asp	Pro 330
Gln	Lys	Tyr	Glu	Ala 335	His	Val	Pro	Glu	Asn 340	Ala	Val	Gly	His	Glu 345
Val	Gln	Arg	Leu	Thr 350	Val	Thr	Asp	Leu	Asp 355	Ala	Pro	Asn	Ser	Pro 360
Ala	Trp	Arg	Ala	Thr 365	Tyr	Leu	Ile	Met	Gly 370	Gly	Asp	Asp	Gly	Asp 375
His	Phe	Thr	Ile	Thr 380	Thr	His	Pro	Glu	Ser 385	Asn	Gln	Gly	Ile	Leu 390
Thr	Thr	Arg	Lys	Gly 395	Leu	Asp	Phe	Glu	Ala 400	Lys	Asn	Gln	His	Thr 405
Leu	Tyr	Val	Glu	Val	Thr	Asn	Glu	Ala	Pro	Phe	Val	Leu	Lys	Leu

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Pro	Thr	Ser	Thr	Ala 425	Thr	Ile	Val	Val	His 430	Val	Glu	Asp	Val	Asn 435
Glu	Ala	Pro	Val	Phe 440	Val	Pro	Pro	Ser	Lys 445	Val	Val	Glu	Val	Gln 450
Glu	Gly	Ile	Pro	Thr 455	Gly	Glu	Pro	Val	Cys 460	Val	Tyr	Thr	Ala	Glu 465
Asp	Pro	Asp	Lys	Glu 470	Asn	Gln	Lys	Ile	Ser 475	Tyr	Arg	Ile	Leu	Arg 480
Asp	Pro	Ala	Gly	Trp 485	Leu	Ala	Met	Asp	Pro 490	Asp	Ser	Gly	Gln	Val 495
Thr	Ala	Val	Gly	Thr 500	Leu	Asp	Arg	Glu	Asp 505	Glu	Gln	Phe	Val	Arg 510
Asn	Asn	Ile	Tyr	Glu 515	Val	Met	Val	Leu	Ala 520	Met	Asp	Asn	Gly	Ser 525
Pro	Pro	Thr	Thr	Gly 530	Thr	Gly	Thr	Leu	Leu 535	Leu	Thr	Leu	Ile	Asp 540
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Cys	Asn	Gln	Ser	Pro 560	Val	Arg	Gln	Val	Leu 565	Asn	Ile	Thr	Asp	Lys 570
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Gly	Ala	Val	Leu	Ala 665	Leu	Leu	Phe	Leu	Leu 670	Leu	Val	Leu	Leu	Leu 675
Leu	Val	Arg	Lys	Lys 680	Arg	Lys	Ile	Lys	Glu 685	Pro	Leu	Leu	Leu	Pro 690
Glu	Asp	Asp	Thr	Arg 695	Asp	Asn	Val	Phe	Tyr 700	Tyr	Gly	Glu	Glu	Gly 705
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Gly Leu Glu Ala Arg Pro Glu Val Val Leu Arg Asn Asp Val Ala 725 730 735 Pro Thr Ile Ile Pro Thr Pro Met Tyr Arg Pro Arg Pro Ala Asn 740 745 Pro Asp Glu Ile Gly Asn Phe Ile Ile Glu Asn Leu Lys Ala Ala 755 760 Asn Thr Asp Pro Thr Ala Pro Pro Tyr Asp Thr Leu Leu Val Phe Asp Tyr Glu Gly Ser Gly Ser Asp Ala Ala Ser Leu Ser Ser Leu 785 790 795 Thr Ser Ser Ala Ser Asp Gln Asp Gln Asp Tyr Asp Tyr Leu Asn Glu Trp Gly Ser Arg Phe Lys Lys Leu Ala Asp Met Tyr Gly Gly 815 820 825

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<212> DNA

<213> Homo sapien

<400> 87

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<211> 1018

<212> PRT

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<400> 88

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Gly Val Ser Glu Glu Asp Lys Gly Phe Gly Pro Ile Phe Glu Glu 35 40 45

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Val	Ser	Leu	Asn	Cys 65	Arg	Ala	Arg	Ala	Ser 70	Pro	Phe	Pro	Val	Tyr 75
Lys	Trp	Arg	Met	Asn 80	Asn	Gly	Asp	Val	Asp 85	Leu	Thr	Ser	Asp	Arg 90
Tyr	Ser	Met	Val	Gly 95	Gly	Asn	Leu	Val	Ile 100	Asn	Asn	Pro	Asp	Lys 105
Gln	Lys	Asp	Ala	Gly 110	Ile	Tyr	Tyr	Cys	Leu 115	Ala	Ser	Asn	Asn	Tyr 120
Gly	Met	Val	Arg	Ser 125	Thr	Glu	Ala	Thr	Leu 130	Ser	Phe	Gly	Tyr	Leu 135
Asp	Pro	Phe	Pro	Pro 140	Glu	Glu	Arg	Pro	Glu 145	Val	Arg	Val	Lys	Glu 150
Gly	Lys	Gly	Met	Val 155	Leu	Leu	Cys	Asp	Pro 160	Pro	Tyr	His	Phe	Pro 165
Asp	Asp	Leu	Ser	Tyr 170	Arg	Trp	Leu	Leu	Asn 175	Glu	Phe	Pro	Val	Phe 180
Ile	Thr	Met	Asp	Lys 185	Arg	Arg	Phe	Val	Ser 190	Gln	Thr	Asn	Gly	Asn 195
Leu	Tyr	Ile	Ala	Asn 200	Val	Glu	Ala	Ser	Asp 205	Lys	Gly	Asn	Tyr	Ser 210
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Cys	Phe	Val	Ser	200 Ser 215	Pro	Ser	Ile	Thr	205 Lys 220	Ser	Val	Phe	_	210 Lys 225
Cys Phe	Phe	Val .Pro	Ser Leu	200 Ser 215 Ile 230	Pro	Ser Ile	Ile Pro	Thr Glu	205 Lys 220 Arg 235	Ser	Val Thr	Phe Lys	Ser	210 Lys 225 Tyr 240
Cys Phe Pro	Phe Ile Ala	Val .Pro Asp	Ser Leu Ile	200 Ser 215 Ile 230 Val 245	Pro Pro Val	Ser Ile Gln	Ile Pro Phe	Thr Glu Lys	205 Lys 220 Arg 235 Asp 250	Ser Thr	Val Thr Tyr	Phe Lys Ala	Ser	210 Lys 225 Tyr 240 Met 255
Cys Phe Pro Gly	Phe Ile Ala	Val .Pro Asp	Ser Leu Ile Val	200 Ser 215 Ile 230 Val 245 Thr 260	Pro Pro Val Leu	Ser Ile Gln Glu	Ile Pro Phe Cys	Thr Glu Lys Phe	205 Lys 220 Arg 235 Asp 250 Ala 265	Ser Thr Val Leu	Val Thr Tyr Gly	Phe Lys Ala Asn	Ser Pro Leu Pro	Lys 225 Tyr 240 Met 255 Val
Cys Phe Pro Gly Pro	Phe Ile Ala Gln	Val .Pro Asp Asn	Ser Leu Ile Val	200 Ser 215 Ile 230 Val 245 Thr 260 Trp 275	Pro Pro Val Leu Arg	Ser Ile Gln Glu Lys	Ile Pro Phe Cys Val	Thr Glu Lys Phe Leu	205 Lys 220 Arg 235 Asp 250 Ala 265 Glu 280	Ser Thr Val Leu Pro	Val Thr Tyr Gly Met	Phe Lys Ala Asn	Ser Pro Leu Pro	210 Lys 225 Tyr 240 Met 255 Val 270
Cys Phe Pro Gly Pro	Phe Ile Ala Gln Asp	Val .Pro Asp Asn Ile	Ser Leu Ile Val Arg	200 Ser 215 Ile 230 Val 245 Thr 260 Trp 275 Thr 290	Pro Pro Val Leu Arg	Ser Ile Gln Glu Lys	Ile Pro Phe Cys Val	Thr Glu Lys Phe Leu Val	205 Lys 220 Arg 235 Asp 250 Ala 265 Glu 280 Leu 295	Ser Thr Val Leu Pro	Val Thr Tyr Gly Met	Phe Lys Ala Asn Pro	Ser Pro Leu Pro Ser Asn	210 Lys 225 Tyr 240 Met 255 Val 270 Thr 285 Ile
Cys Phe Pro Gly Pro Ala Gln	Phe Ile Ala Gln Asp Glu Leu	Val Pro Asp Asn Ile Ile	Ser Leu Ile Val Arg Ser Asp	200 Ser 215 Ile 230 Val 245 Thr 260 Trp 275 Thr 290 Glu 305	Pro Pro Val Leu Arg Ser Gly	Ser Ile Gln Glu Lys Gly Ile	Ile Pro Phe Cys Val Ala	Thr Glu Lys Phe Leu Val Glu	205 Lys 220 Arg 235 Asp 250 Ala 265 Glu 280 Leu 295 Cys 310	Ser Thr Val Leu Pro Lys Glu	Val Thr Tyr Gly Met Ile Ala	Phe Lys Ala Asn Pro Phe Glu	Ser Pro Leu Pro Ser Asn	210  Lys 225  Tyr 240  Met 255  Val 270  Thr 285  Ile 300  Ile 315

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Thr	Ile	Arg	Trp	Leu 365	Lys	Asn	Gly	Tyr	Ala 370	Tyr	His	Lys	Gly	Glu 375
Leu	Arg	Leu	Tyr	Asp 380	Val	Thr	Phe	Glu	Asn 385	Ala	Gly	Met	Tyr	Gln 390
Cys	Ile	Ala	Glu	Asn 395	Thr	Tyr	Gly	Ala	Ile 400	Tyr	Ala	Asn	Ala	Glu 405
Leu	Lys	Ile	Leu	Ala 410	Leu	Ala ,	Pro	Thr	Phe 415	Glu	Met	Asn	Pro	Met 420
Lys	Lys	Lys	Ile	Leu 425	Ala	Ala	Lys	Gly	Gly 430	Arg	Val	Ile	Ile	Glu 435
Cys	Lys	Pro	Lys	Ala 440	Ala	Pro	Lys	Pro	Lys 445	Phe	Ser	Trp	Ser	Lys 450
Gly	Thr	Glu	Trp	Leu 455	Val	Asn	Ser	Ser	Arg 460	Ile	Leu	Ile	Trp	Glu 465
Asp	Gly	Ser	Leu	Glu 470	Ile	Asn	Asn	Ile	Thr 475	Arg	Asn	Asp	Gly	Gly 480
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Thr	Gly	Thr	Leu	Val 500	Ile	Thr	Asp	Pro	Thr 505	Arg	Ile	Ile	Leu	Ala 510
Pro	Ile	Asn	Ala	Asp 515	Ile	Thr	Val	Gly	Glu 520	Asn	Ala	Thr	Met	Gln 525
Cys	Ala	Ala	Ser	Phe 530	Asp	Pro	Ala	Leu	Asp 535	Leu	Thr	Phe	Val	Trp 540
Ser	Phe	Asn	Gly	Tyr 545	Val	Ile	Asp	Phe	Asn 550	Lys	Glu	Asn	Ile	His 555
Tyr	Gln	Arg	Asn	Phe 560	Met	Leu	Asp	Ser	Asn 565	Gly	Glu	Leu	Leu	Ile 570
Arg	Asn	Ala	Gln	Leu 575	Lys	His	Ala	Gly	Arg 580	Tyr	Thr	Cys	Thr	Ala 585
Gln	Thr	Ile	Val	Asp 590	Asn	Ser	Ser	Ala	Ser 595	Ala	Asp	Leu	Val	Val 600
Arg	Gly	Pro	Pro	Gly 605	Pro	Pro	Gly	Gly	Leu 610	Arg	Ile	Glu	Asp	Ile 615
Arg	Ala	Thr	Ser	Val 620	Ala	Leu	Thr	Trp	Ser 625	Arg	Gly	Ser	Asp	Asn 630
His	Ser	Pro	Ile	Ser 635	Lys	Tyr	Thr	Ile	Gln 640	Thr	Lys	Thr	Ile	Leu 645
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Glu	Pro	Ser	Ile	Pro 695	Ser	Asn	Arg	Ile	Lys 700	Thr	Asp	Gly	Ala	Ala 705
Pro	Asn	Val	Ala	Pro 710	Ser	Asp	Val	Gly	Gly 715	Gly	Gly	Gly	Arg	Asn 720
Arg	Glu	Leu	Thr	Ile 725	Thr	Trp	Ala	Pro	Leu 730	Ser	Arg	Glu	Tyr	His 735
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Ser	Ala	Gly	Cys	Gly 890	Pro	Pro	Ser	Asp	Met 895	Ile	Glu	Ala	Phe	Thr 900
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Arg	Ser	Gly	Ser	Arg 920	Tyr	Ile	Ile	Thr	Trp 925	Asp	His	Val	Val	Ala 930
Leu	Ser	Asn	Glu	Ser 935	Thr	Val	Thr	Gly	Tyr 940	Lys	Val	Leu	Tyr	Arg 945
Pro	Asp	Gly	Gln	His 950	Asp	Gly	Lys	Leu	Tyr 955	Ser	Thr	His	Lys	His 960

بيديد

Ser Ile Glu Val Pro Ile Pro Arg Asp Gly Glu Tyr Val Val Glu 965 970 975

Val Arg Ala His Ser Asp Gly Gly Asp Gly Val Val Ser Gln Val 980 985 990

Lys Ile Ser Gly Ala Pro Thr Leu Ser Pro Ser Leu Leu Gly Leu 995 1000 1005

Leu Leu Pro Ala Phe Gly Ile Leu Val Tyr Leu Glu Phe  $1010 \\ \hspace{1.5cm} 1015$ 

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<211> 1616

<212> DNA

<213> Homo sapien

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<211> 300

<212> PRT

<213> Homo sapien

<400> 90

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Gln Leu Tyr Asn Lys Tyr Pro Asp Ala Val Ala Thr Trp Leu Asn
45

Pro Asp Pro Ser Gln Lys Gln Asn Leu Leu Ala Pro Gln Thr Leu
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Pro Ser Lys Ser Asn Glu Ser His Asp His Met Asp Asp Met Asp
65 70 75

Asp Glu Asp Asp Asp Asp His Val Asp Ser Gln Asp Ser Ile Asp 80 85 90

Ser Asn Asp Ser Asp Asp Val Asp Asp Thr Asp Asp Ser His Gln 95 100 105

Ser Asp Glu Ser His His Ser Asp Glu Ser Asp Glu Leu Val Thr 110 115 120

Asp Phe Pro Thr Asp Leu Pro Ala Thr Glu Val Phe Thr Pro Val 125 130 130

Val Pro Thr Val Asp Thr Tyr Asp Gly Arg Gly Asp Ser Val Val 140 145 150

Tyr Gly Leu Arg Ser Lys Ser Lys Phe Arg Arg Pro Asp Ile

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Gln	Tyr	Pro	Asp	Ala 170	Thr	Asp	Glu	Asp	Ile 175	Thr	Ser	His	Met	Glu 180
Ser	Glu	Glu	Leu	Asn 185	Gly	Ala	Tyr	Lys	Ala 190	Ile	Pro	Val	Ala	Gln 195
Asp	Leu	Asn	Ala	Pro 200	Ser	Asp	Trp	Asp	Ser 205	Arg	Gly	Lys	Asp	Ser 210
Tyr	Glu	Thr	Ser	Gln 215	Leu	Asp	Asp	Gln	Ser 220	Ala	Glu	Thr	His	Ser 225
His	Lys	Gln	Ser	Arg 230	Leu	Tyr	Lys	Arg	Lys 235	Ala	Asn	Asp	Glu	Ser 240
Asn	Glu	His	Ser	Asp 245	Val	Ile	Asp	Ser	Gln 250	Glu	Leu	Ser	Lys	Val 255
Ser	Arg	Glu	Phe	His 260	Ser	His	Glu	Phe	His 265	Ser	His	Glu	Asp	Met 270
Leu	Val	Val	Asp	Pro 275	Lys	Ser	Lys	Glu	Glu 280	Asp	Lys	His	Leu	Lys 285
Phe	Arg	Ile	Ser	His 290	Glu	Leu	Asp	Ser	Ala 295	Ser	Ser	Glu	Val	Asn 300
<210	> 91													
<2112	> 259	93												
<212	> DNA	A												

<213> Homo sapien

## <400> 91

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atcacctatg acacgccttt caaaagagaa aagtcttttg agatcgtgat 700 tatggtgcta aaggacaaat tccaggtggc tgtaaatgga aaacatactc 750 tgctctatgg ccacaggatc ggcccagaga aaatagacac tctgggcatt 800 tatggcaaag tgaatattca ctcaattggt tttagcttca gctcggactt 850 acaaagtacc caagcatcta gtctggaact gacagagata agtagagaaa 900 atgttccaaa gtctggcacg ccccagcttc agactgtctc tccctcctgg 950 gatttacagg gtcatggctc tgaaacattc tgtagtgttc tttggacacg 1000 agttttcctg gagatcgctt tctgcaggcc tattggtctg actgtggctt 1050 cttttcagag cctgccattc gctgcaaggt tgaacacccc catgggccct 1100 ggacgaactg tcgtcgttaa aggagaagtg aatgcaaatg ccaaaagctt 1150 taatgttgac ctactagcag gaaaatcaaa ggatattgct ctacacttga 1200 acccacgcct gaatattaaa gcatttgtaa gaaattcttt tcttcaggag 1250 teetggggag aagaagagag aaatattace tettteeeat ttagteetgg 1300 gatgtacttt gagatgataa tttactgtga tgttagagaa ttcaaggttg 1350 cagtaaatgg cgtacacagc ctggagtaca aacacagatt taaagagctc 1400 agcagtattg acacgctgga aattaatgga gacatccact tactggaagt 1450 aaggagetgg tageetacet acacagetge tacaaaaace aaaatacaga 1500 atggcttctg tgatactggc cttgctgaaa cgcatctcac tgtcattcta 1550 ttgtttatat tgttaaaatg agcttgtgca ccattagatc ctgctgggtg 1600 ttctcagtcc ttgccatgaa gtatggtggt gtctagcact gaatggggaa 1650 actgggggca gcaacactta tagccagtta aagccactct gccctctctc 1700 ctactttggc tgactcttca agaatgccat tcaacaagta tttatggagt 1750 acctactata atacagtage taacatgtat tgageacaga ttttttttgg 1800 taaaactgtg aggagctagg atatatactt ggtgaaacaa accagtatgt 1850 tccctgttct cttgagcttc gactcttctg tgctctattg ctgcgcactg 1900 ctttttctac aggcattaca tcaactccta aggggtcctc tgggattagt 1950 taagcagcta ttaaatcacc cgaagacact aatttacaga agacacaact 2000 ccttccccag tgatcactgt cataaccagt gctctaccgt atcccatcac 2050 tgaggactga tgttgactga catcatttta tcgtaataaa catgtggctc 2100 tattagctgc aagctttacc aagtaattgg catgacatct gagcacagaa 2150 attaaggcaa aaaaccaaag caaaacaaat acatggtgct gaaattaact 2200

<210> 92

<211> 359

<212> PRT

<213> Homo sapien

<400> 92

Met Leu Ser Leu Asn Asn Leu Gln Asn Ile Ile Tyr Asn Pro Val 1 5 10 15

Ile Pro Tyr Val Gly Thr Ile Pro Asp Gln Leu Asp Pro Gly Thr 20 25 30

Leu Ile Val Ile Cys Gly His Val Pro Ser Asp Ala Asp Arg Phe 35 40 45

Gln Val Asp Leu Gln Asn Gly Ser Ser Val Lys Pro Arg Ala Asp
50 55 60

Val Ala Phe His Phe Asn Pro Arg Phe Lys Arg Ala Gly Cys Ile 65 70 75

Val Cys Asn Thr Leu Ile Asn Glu Lys Trp Gly Arg Glu Glu Ile 80 85 90 41.7 ×

Thr Tyr Asp Thr Pro Phe Lys Arg Glu Lys Ser Phe Glu Ile Val 95 100 105

Ile Met Val Leu Lys Asp Lys Phe Gln Val Ala Val Asn Gly Lys
110 115 120

His Thr Leu Leu Tyr Gly His Arg Ile Gly Pro Glu Lys Ile Asp 125 130 135

Thr Leu Gly Ile Tyr Gly Lys Val Asn Ile His Ser Ile Gly Phe 140 145 150

Ser Phe Ser Ser Asp Leu Gln Ser Thr Gln Ala Ser Ser Leu Glu 155 160 165

Leu Thr Glu Ile Ser Arg Glu Asn Val Pro Lys Ser Gly Thr Pro 170 175 180

Gln Leu Gln Thr Val Ser Pro Ser Trp Asp Leu Gln Gly His Gly 185 190 195

Ser Glu Thr Phe Cys Ser Val Leu Trp Thr Arg Val Phe Leu Glu 200 205 210 Ile Ala Phe Cys Arg Pro Ile Gly Leu Thr Val Ala Ser Phe Gln 215 220 Ser Leu Pro Phe Ala Ala Arg Leu Asn Thr Pro Met Gly Pro Gly 230 235 240 Arg Thr Val Val Lys Gly Glu Val Asn Ala Asn Ala Lys Ser 245 250 Phe Asn Val Asp Leu Leu Ala Gly Lys Ser Lys Asp Ile Ala Leu 260 265 270 His Leu Asn Pro Arg Leu Asn Ile Lys Ala Phe Val Arg Asn Ser 280 Phe Leu Gln Glu Ser Trp Gly Glu Glu Glu Arg Asn Ile Thr Ser 295 300 Phe Pro Phe Ser Pro Gly Met Tyr Phe Glu Met Ile Ile Tyr Cys 310 Asp Val Arg Glu Phe Lys Val Ala Val Asn Gly Val His Ser Leu Glu Tyr Lys His Arg Phe Lys Glu Leu Ser Ser Ile Asp Thr Leu Glu Ile Asn Gly Asp Ile His Leu Leu Glu Val Arg Ser Trp

<210> 93

<211> 2401

<212> DNA

<213> Homo sapien

## <400> 93

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tocaagatoo atgagaaggo ottoagooca otgoggaago tgoagaagot 600 ctacatetee aagaaceace tggtggagat cccgcccaac ctacccaget 650 ccctggtgga gctccgcatc cacgacaacc gcatccgcaa ggtgcccaag 700 ggagtgttca gcgggctccg gaacatgaac tgcatcgaga tgggcgggaa 750 cccactggag aacagtggct ttgaacctgg agccttcgat ggcctgaagc 800 tcaactacct gcgcatctca gaggccaagc tgactggcat ccccaaagac 850 ctccctgaga ccctgaatga actccaccta gaccacaaca aaatccaggc 900 catcgaactg gaggacctgc ttcgctactc caagctgtac aggctgggcc 950 taggccacaa ccagatcagg atgatcgaga acgggagcct gagcttcctg 1000 cccaccetce gggageteca ettggacaac aacaagttgg ccagggtgcc 1050 ctcagggctc ccagacctca agctcctcca ggtggtctat ctgcactcca 1100 acaacatcac caaagtgggt gtcaacgact tctgtcccat gggcttcggg 1150 gtgaagcggg cctactacaa cggcatcagc ctcttcaaca accccgtgcc 1200 ctactgggag gtgcagccgg ccactttccg ctgcgtcact gaccgcctgg 1250 ccatccagtt tggcaactac aaaaagtaga ggcagctgca gccaccgcgg 1300 ggcctcagtg ggggtctctg gggaacacag ccagacatcc tgatggggag 1350 gcagagccag gaagctaagc cagggcccag ctgcgtccaa cccagccccc 1400 cacctegggt ceetgacee agetegatge eccateaceg cetetecetg 1450 gctcccaagg gtgcaggtgg gcgcaaggcc cggcccccat cacatgttcc 1500 ettggcetca gagetgeece tgeteteeca ceacageeae ecagaggeae 1550 cccatgaagc ttttttctcg ttcactccca aacccaagtg tccaaggctc 1600 cagtectagg agaacagtee etgggteage agecaggagg eggteeataa 1650 gaatggggac agtgggctct gccagggctg ccgcacctgt ccagacacac 1700 atgttctgtt cctcctcctc atgcatttcc agcctttcaa ccctccccga 1750 ctctgcggct cccctcagcc cccttgcaag ttcatggcct gtccctccca 1800 gacccctgct ccactggccc ttcgaccagt cctcccttct gttctctctt 1850 teccegteet tectetet etetetet etetetet ettetetet 1900 gtgtgtgtgt gtgtgtgtt gtgtgtgtgt gtgtgtgtgt cttgtgcttc 1950 ctcagacctt tctcgcttct gagcttggtg gcctgttccc tccatctctc 2000 cgaacctggc ttcgcctgtc cctttcactc cacaccctct ggccttctgc 2050

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<210> 94

<211> 368

<212> PRT

<213> Homo sapien

<400> 94

Met Trp Pro Leu Trp Arg Leu Val Ser Leu Leu Ala Leu Ser Gln 1 5 10 15

Ala Leu Pro Phe Glu Gln Arg Gly Phe Trp Asp Phe Thr Leu Asp
20 25 30

Asp Gly Pro Phe Met Met Asn Asp Glu Glu Ala Ser Gly Ala Asp 35 40 45

Thr Ser Gly Val Leu Asp Pro Asp Ser Val Thr Pro Thr Tyr Ser 50 55 60

Ala Met Cys Pro Phe Gly Cys His Cys His Leu Arg Val Val Gln
65 70 75

Cys Ser Asp Leu Gly Leu Lys Ser Val Pro Lys Glu Ile Ser Pro 80 85 90

e ..

Asp Thr Thr Leu Leu Asp Leu Gln Asn Asn Asp Ile Ser Glu Leu 95 100 105

Arg Lys Asp Asp Phe Lys Gly Leu Gln His Leu Tyr Ala Leu Val 110 115 120

Leu Val Asn Asn Lys Ile Ser Lys Ile His Glu Lys Ala Phe Ser 125 130 135

Pro Leu Arg Lys Leu Gln Lys Leu Tyr Ile Ser Lys Asn His Leu 140 145 150

Val Glu Ile Pro Pro Asn Leu Pro Ser Ser Leu Val Glu Leu Arg
155 160 165

Ile His Asp Asn Arg Ile Arg Lys Val Pro Lys Gly Val Phe Ser 170 175 180

Gly Leu Arg Asn Met Asn Cys Ile Glu Met Gly Gly Asn Pro Leu 185 190 195

Glu Asn Ser Gly Phe Glu Pro Gly Ala Phe Asp Gly Leu Lys Leu 200 205 210 Asn Tyr Leu Arg Ile Ser Glu Ala Lys Leu Thr Gly Ile Pro Lys 215 220 225 Asp Leu Pro Glu Thr Leu Asn Glu Leu His Leu Asp His Asn Lys 230 235 Ile Gln Ala Ile Glu Leu Glu Asp Leu Leu Arg Tyr Ser Lys Leu 250 255 Tyr Arg Leu Gly Leu Gly His Asn Gln Ile Arg Met Ile Glu Asn 265 Gly Ser Leu Ser Phe Leu Pro Thr Leu Arg Glu Leu His Leu Asp 275 280 285 Asn Asn Lys Leu Ala Arg Val Pro Ser Gly Leu Pro Asp Leu Lys 300 Leu Leu Gln Val Val Tyr Leu His Ser Asn Asn Ile Thr Lys Val 310 315 Gly Val Asn Asp Phe Cys Pro Met Gly Phe Gly Val Lys Arg Ala 325 330 Tyr Tyr Asn Gly Ile Ser Leu Phe Asn Asn Pro Val Pro Tyr Trp 345 Glu Val Gln Pro Ala Thr Phe Arg Cys Val Thr Asp Arg Leu Ala Ile Gln Phe Gly Asn Tyr Lys Lys

<210> 95

<211> 1983

<212> DNA

<213> Homo sapien

## <400> 95

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ctgccccgcc tgctgctgc gctgctgctg ctgcccgccg ccgggccggc 200
ccagttccac ggggagaagg gcatctccat cccggaccac ggcttctgcc 250
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atgcccaacc ttctgggcca cacgaaccag gaggacgcag gcctagaggt 350
gcaccagttc tatccgctgg tgaaggtgca gtgctcgccc gaactgcgct 400
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<210> 96 <211> 565 <212> PRT <213> Homo sapien <400> 96 Met Arg Pro Arg Ser Ala Leu Pro Arg Leu Leu Pro Leu Leu Leu Leu Pro Ala Ala Gly Pro Ala Gln Phe His Gly Glu Lys Gly Ile Ser Ile Pro Asp His Gly Phe Cys Gln Pro Ile Ser Ile Pro Leu Cys Thr Asp Ile Ala Tyr Asn Gln Thr Ile Met Pro Asn Leu Leu Gly His Thr Asn Gln Glu Asp Ala Gly Leu Glu Val His Gln Phe Tyr Pro Leu Val Lys Val Gln Cys Ser Pro Glu Leu Arg Phe Phe Leu Cys Ser Met Tyr Ala Pro Val Cys Thr Val Leu Glu Gln Ala Ile Pro Pro Cys Arg Ser Ile Cys Glu Arg Ala Arg Gln Gly 110 115 Cys Glu Ala Leu Met Asn Lys Phe Gly Phe Gln Trp Pro Glu Arg 125 Leu Arg Cys Glu His Phe Pro Arg His Gly Ala Glu Gln Ile Cys Val Gly Gln Asn His Ser Glu Asp Gly Ala Pro Ala Leu Leu Thr Thr Ala Pro Pro Pro Gly Leu Gln Pro Gly Ala Gly Gly Thr Pro Gly Gly Pro Gly Gly Gly Ala Pro Pro Arg Tyr Ala Thr Leu 195 Glu His Pro Phe His Cys Pro Arg Val Leu Lys Val Pro Ser Tyr 205 Leu Ser Tyr Lys Phe Leu Gly Glu Arg Asp Cys Ala Ala Pro Cys 215 220 225 Glu Pro Ala Arg Pro Asp Gly Ser Met Phe Phe Ser Gln Glu Glu 235 Thr Arg Phe Ala Arg Leu Trp Ile Leu Thr Trp Ser Val Leu Cys 245 250 Cys Ala Ser Thr Phe Phe Thr Val Thr Tyr Leu Val Asp Met

Gln Arg Phe Arg Tyr Pro Glu Arg Pro Ile Ile Phe Leu Ser Gly

				275					280					285
Cys	Tyr	Thr	Met	Val 290	Ser	Val	Ala	Tyr	Ile 295	Ala	Gly	Phe	Val	Leu 300
Gln	Glu	Arg	Val	Val 305	Суѕ	Asn	Glu	Arg	Phe 310	Ser	Glu	Asp	Gly	Tyr 315
Arg	Thr	Val	Val	Gln 320	Gly	Thr	Lys	Lys	Glu 325	Gly	Cys	Thr	Ile	Leu 330
Phe	Met	Met	Leu	Tyr 335	Phe	Phe	Ser	Met	Ala 340	Ser	Ser	Ile	Trp	Trp 345
Val	Ile	Leu	Ser	Leu 350	Thr	Trp	Phe	Leu	Ala 355	Ala	Gly	Met	Lys	Trp 360
Gly	His	Glu	Ala	Ile 365	Glu	Ala	Asn	Ser	Gln 370	Tyr	Phe	His	Leu	Ala 375
Ala	Trp	Ala	Val	Pro 380	Ala	Val	Lys	Thr	Ile 385	Thr	Ile	Leu	Ala	Met 390
Gly	Gln	Ile	Asp	Gly 395	Asp	Leu	Leu	Ser	Gly 400	Val	Cys	Phe	Val	Gly 405
Leu	Asn	Ser	Leu	Asp 410	Pro	Leu	Arg	Gly	Phe 415	Val	Leu	Ala	Pro	Leu 420
Phe	Val	Tyr	Leu	Phe 425	Ile	Gly	Thr	Ser	Phe 430	Leu	Leu	Ala	Gly	Phe 435
Val	Ser	Leu	Phe	Arg 440	Ile	Arg	Thr	Ile	Met 445	Lys	His	Asp	Gly	Thr 450
Lys	Thr	Glu	Lys	Leu 455	Glu	Arg	Leu	Met	Val 460	Arg	Ile	Gly	Val	Phe 465
Ser	Val	Leu	Tyr	Thr 470	Val	Pro	Ala	Thr	Ile 475	Val	Ile	Ala	Cys	Tyr 480
Phe	Tyr	Glu	Gln	Ala 485	Phe	Arg	Glu	His	Trp 490	Glu	Arg	Ser	Trp	Val 495
Ser	Gln	His	Cys	Lys 500	Ser	Leu	Ala	Ile	Pro 505	Cys	Pro	Ala	His	Tyr 510
Thr	Pro	Arg	Met	Ser 515	Pro	Asp	Phe	Thr	Val 520	Tyr	Met	Ile	Lys	Tyr 525
Leu	Met	Thr	Leu	Ile 530	Val	Gly	Ile	Thr	Ser 535	Gly	Phe	Trp	Ile	Trp 540
Ser	Gly	Lys	Thr	Leu 545	His	Ser	Trp	Arg	Lys 550	Phe	Tyr	Thr	Arg	Leu 555
Thr	Asn	Ser	Arg	His 560	Gly	Glu	Thr	Thr	Val 565					

<210> 97 <211> 2294 <212> DNA <213> Homo sapien

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<210> 98

<211> 428

<212> PRT

<213> Homo sapien

## <400> 98

Met Gln Glu Leu His Leu Leu Trp Trp Ala Leu Leu Leu Gly Leu 1 5 10 15

Ala Gln Ala Cys Pro Glu Pro Cys Asp Cys Gly Glu Lys Tyr Gly
20 25 30

Phe Gln Ile Ala Asp Cys Ala Tyr Arg Asp Leu Glu Ser Val Pro 35 40 45

Pro Gly Phe Pro Ala Asn Val Thr Thr Leu Ser Leu Ser Ala Asn 50 55 60

Arg Leu Pro Gly Leu Pro Glu Gly Ala Phe Arg Glu Val Pro Leu 65 70 75

Leu Gln Ser Leu Trp Leu Ala His Asn Glu Ile Arg Thr Val Ala  $80 \\ 85 \\ 90$ 

Ala Gly Ala Leu Ala Ser Leu Ser His Leu Lys Ser Leu Asp Leu 95 100 105

Ser	His	Asn	Leu	Ile 110	Ser	Asp	Phe	Ala	Trp 115	Ser	Asp	Leu	His	Asn 120
Leu	Ser	Ala	Leu	Gln 125	Leu	Leu	Lys	Met	Asp 130	Ser	Asn	Glu	Leu	Thr 135
Phe	Ile	Pro	Arg	Asp 140	Ala	Phe	Arg	Ser	Leu 145	Arg	Ala	Leu	Arg	Ser 150
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Gly Gln Asp Asn	Thr Pro Phe Glu Pro	Asp Arg Gly Leu Gly	Gly Phe Lys Gly	275 Arg 290 Gln 305 Leu 320 Ser 335 Gly 350	Ala Ala Glu Ala	Leu Phe Glu Glu	Pro Ala Gly Ser Asp	Gly Asn Thr Ser	280 Thr 295 Gly 310 Tyr 325 Val 340 Leu 355	Pro Ser Ser Asp	Val Leu Cys Val Arg	Ala Leu Leu Ala Arg	Ser Ile Ala Leu Phe	285 Ser 300 Pro 315 Thr 330 Ala 345 His 360
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Ile	Thr	Arg	Glu	Asp 185	Gln	Tyr	Ser	Lys	Pro 190	Pro	His	Lys	Lys	Leu 195
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Ala Gly Glu Gly Pro Phe Ser Glu Thr Tyr Thr Phe Ser Thr Thr 1040 1045 1050

Lys Ser Val Pro Pro Thr Ile Lys Ala Pro Arg Val Thr Gln Leu 1055 1060 1065

Glu Gly Asn Ser Cys Glu Ile Leu Trp Glu Thr Val Pro Ser Met 1070 1075 1080

Lys Gly Asp Pro Val Asn Tyr Ile Leu Gln Val Leu Val Gly Arg 1085 1090 1095

Glu Ser Glu Tyr Lys Gln Val Tyr Lys Gly Glu Glu Ala Thr Phe 1100 1105 1110

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Cys Ala Cys Arg Arg Cys Leu Asp Thr Ser Gln Glu Leu Ser Gly
1130 1135 1140

Ala Phe Ser Pro Ser Ala Ala Phe Val Leu Gln Arg Ser Glu Val 1145 1150 1155

Met Leu Thr Gly Asp Met Gly Ser Leu Asp Asp Pro Lys Met Lys 1160 1165 1170

Ser Met Met Pro Thr Asp Glu Gln Phe Ala Ala Ile Ile Val Leu 1175 1180 1180

Gly Phe Ala Thr Leu Ser Ile Leu Phe Ala Phe Ile Leu Gln Tyr 1190 1195 1200

Phe Leu Met Lys

<210> 101

<211> 2565

<212> DNA

<213> Homo sapien

<400> 101

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<210> 102

<211> 757

<212> PRT

<213> Homo sapien

<400> 102

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20 25 30

Pro Ser Ser Cys Tyr Ala Leu Phe Pro Arg Arg Thr Phe Leu
35 40 45

Glu Ala Trp Arg Ala Cys Arg Glu Leu Gly Gly Asp Leu Ala Thr
50 55 60

Pro Arg Thr Pro Glu Glu Ala Gln Arg Val Asp Ser Leu Val Gly 65 70 75

Ala Gly Pro Ala Ser Arg Leu Leu Trp Ile Gly Leu Gln Arg Gln
80 85 90

Ala	Arg	Gln	Cys	Gln 95	Leu	Gln	Arg	Pro	Leu 100	Arg	Gly	Phe	Thr	Trp 105
Thr	Thr	Gly	Asp	Gln 110	Asp	·Thr	Ala	Phe	Thr 115	Asn	Trp	Ala	Gln	Pro 120
Ala	Ser	Gly	Gly	Pro 125	Cys	Pro	Ala	Gln	Arg 130	Cys	Val	Ala	Leu	Glu 135
Ala	Ser	Gly	Glu	His 140	Arg	Trp	Leu	Glu	Gly 145	Ser	Cys	Thr	Leu	Ala 150
Val	Asp	Gly	Tyr	Leu 155	Cys	Gln	Phe	Gly	Phe 160	Glu	Gly	Ala	Cys	Pro 165
Ala	Leu	Gln	Asp	Glu 170	Ala	Gly	Gln	Ala	Gly 175	Pro	Ala	Val	Tyr	Thr 180
Thr	Pro	Phe	His	Leu 185	Val	Ser	Thr	Glu	Phe 190	Glu	Trp	Leu	Pro	Phe 195
Gly	Ser	Val	Ala	Ala 200	Val	Gln	Cys	Gln	Ala 205	Gly	Arg	Gly	Ala	Ser 210
Leu	Leu	Cys	Val	Lys 215	Gln	Pro	Glu	Gly	Gly 220	Val	Gly	Trp	Ser	Arg 225
Ala	Gly	Pro	Leu	Cys 230	Leu	Gly	Thr	Gly	Cys 235	Ser	Pro	Asp	Asn	Gly 240
Gly	Cys	Glu	His	Glu 245	Cys	Val	Glu	Glu	Val 250	Asp	Gly	His	Val	Ser 255
Cys	Arg	Cys	Thr	Glu 260	Gly	Phe	Arg	Leu	Ala 265	Ala	Asp	Gly	Arg	Ser 270
Cys	Glu	Asp	Pro	Cys 275	Ala	Gln	Ala	Pro	Cys 280	Glu	Gln	Gln	Cys	Glu 285
Pro	Gly	Gly	Pro	Gln 290	Gly	Tyr	Ser	Cys	His 295	Cys	Arg	Leu	Gly	Phe 300
Arg	Pro	Ala	Glu	Asp 305	Asp	Pro	His	Arg	Cys 310	Val	Asp	Thr	Asp	Glu 315
Cys	Gln	Ile	Ala	Gly 320	Val	Cys	Gln	Gln	Met 325	Cys	Val	Asn	Tyr	Val 330
Gly	Gly	Phe	Glu	Cys 335	Tyr	Cys	Ser	Glu	Gly 340	His	Glu	Leu	Glu	Ala 345
Asp	Gly	Ile	Ser	Cys 350	Ser	Pro	Ala	Gly	Ala 355	Met	Gly	Ala	Gln	Ala 360
Ser	Gln	Asp	Leu	Gly 365	Asp	Glu	Leu	Leu	Asp 370	Asp	Gly	Glu	Asp	Glu 375
Glu	Asp	Glu	Asp	Glu 380	Ala	Trp	Lys	Ala	Phe 385	Asn	Gly	Gly	Trp	Thr 390
Glu	Met	Pro	Gly	Ile	Leu	Trp	Met	Glu	Pro	Thr	Gln	Pro	Pro	Asp

				395					400					405
Phe	Ala	Leu	Ala	Tyr 410	Arg	Pro	Ser	Phe	Pro 415	Glu	Asp	Arg	Glu	Pro 420
Gln	Ile	Pro	Tyr	Pro 425	Glu	Pro	Thr	Trp	Pro 430	Pro	Pro	Leu	Ser	Ala 435
Pro	Arg	Val	Pro	Tyr 440	His	Ser	Ser	Val	Leu 445	Ser	Val	Thr	Arg	Pro 450
Val	Val	Val	Ser	Ala 455	Thr	His	Pro	Thr	Leu 460	Pro	Ser	Ala	His	Gln 465
Pro	Pro	Val	Ile	Pro 470	Ala	Thr	His	Pro	Ala 475	Leu	Ser	Arg	Asp	His 480
Gln	Ile	Pro	Val	Ile 485	Ala	Ala	Asn	Tyr	Pro 490	Asp	Leu	Pro	Ser	Ala 495
Tyr	Gln	Pro	Gly	Ile 500	Leu	Ser	Val	Ser	His 505	Ser	Ala	Gln	Pro	Pro 510
Ala	His	Gln	Pro	Pro 515	Met	Ile	Ser	Thr	Lys 520	Tyr	Pro	Glu	Leu	Phe 525
Pro	Ala	His	Gln	Ser 530	Pro	Met	Phe	Pro	Asp 535	Thr	Arg	Val	Ala	Gly 540
Thr	Gln	Thr	Thr	Thr 545	His	Leu	Pro	Gly	Ile 550	Pro	Pro	Asn	His	Ala 555
Pro	Leu	Val	Thr	Thr 560	Leu	Gly	Ala	Gln	Leu 565	Pro	Pro	Gln	Ala	Pro 570
Asp	Ala	Leu	Val	Leu 575	Arg	Thr	Gln	Ala	Thr 580	Gln	Leu	Pro	Ile	Ile 585
Pro	Thr	Ala	Gln	Pro 590	Ser	Leu	Thr	Thr	Thr 595	Ser	Arg	Ser	Pro	Val 600
Ser	Pro	Ala	His	Gln 605	Ile	Ser	Val	Pro	Ala 610	Ala	Thr	Gln	Pro	Ala 615
Ala	Leu	Pro	Thr	Leu 620	Leu	Pro	Ser	Gln	Ser 625	Pro	Thr	Asn	Gln	Thr 630
Ser	Pro	Ile	Ser	Pro 635	Thr	His	Pro	His	Ser 640	Lys	Ala	Pro	Gln	Ile 645
Pro	Arg	Glu	Asp	Gly 650	Pro	Ser	Pro	Lys	Leu 655	Ala	Leu	Trp	Leu	Pro 660
Ser	Pro	Ala	Pro	Thr 665	Ala	Ala	Pro	Thr	Ala 670	Leu	Gly	Glu	Ala	Gly 675
Leu	Ala	Glu	His	Ser 680	Gln	Arg	Asp	Asp	Arg 685	Trp	Leu	Leu	Val	Ala 690
Leu	Leu	Val	Pro	Thr 695	Cys	Val	Phe	Leu	Val 700	Val	Leu	Leu	Ala	Leu 705

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Gly Ile Val Tyr Cys Thr Arg Cys Gly Pro His Ala Pro Asn Lys 710 715 720

Arg Ile Thr Asp Cys Tyr Arg Trp Val Ile His Ala Gly Ser Lys 725 730 735

Ser Pro Thr Glu Pro Met Pro Pro Arg Gly Ser Leu Thr Gly Val 740  $\phantom{000}745$   $\phantom{000}745$ 

Gln Thr Cys Arg Thr Ser Val 755

<210> 103

<211> 2269

<212> DNA

<213> Homo sapien

<400> 103

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<210> 104

<211> 496

<212> PRT

<213> Homo sapien

<400> 104

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Ala Ala Ala Tyr Asn Asn Phe Arg Lys Ser Met Asp Ser Ile Gly
20 25 30

Lys	Lys	Gln	Tyr	Gln 35	Val	Gln	His	Gly	Ser 40	Cys	Ser	Tyr	Thr	Phe 45
Leu	Leu	Pro	Glu	Met 50	Asp	Asn	Cys	Arg	Ser 55	Ser	Ser	Ser	Pro	Tyr 60
Val	Ser	Asn	Ala	Val 65	Gln	Arg	Asp	Ala	Pro 70	Leu	Glu	Tyr	Asp	Asp 75
Ser	Val	Gln	Arg	Leu 80	Gln	Val	Leu	Glu	Asn 85	Ile	Met	Glu	Asn	Asn 90
Thr	Gln	Trp	Leu	Met 95	Lys	Leu	Glu	Asn	Tyr 100	Ile	Gln	Asp	Asn	Met 105
Lys	Lys	Glu	Met	Val 110	Glu	Ile	Gln	Gln	Asn 115	Ala	Val	Gln	Asn	Gln 120
Thr	Ala	Val	Met	Ile 125	Glu	Ile	Gly	Thr	Asn 130	Leu	Leu	Asn	Gln	Thr 135
Ala	Glu	Gln	Thr	Arg 140	Lys	Leu	Thr	Asp	Val 145	Glu	Ala	Gln	Val	Leu 150
Asn	Gln	Thr	Thr	Arg 155	Leu	Glu	Leu	Gln	Leu 160	Leu	Glu	His	Ser	Leu 165
Ser	Thr	Asn	Lys	Leu 170	Glu	Lys	Gln	Ile	Leu 175	Asp	Gln	Thr	Ser	Glu 180
Ile	Asn	Lys	Leu	Gln 185	Asp	Lys	Asn	Ser	Phe 190	Leu	Glu	Lys	Lys	Val 195
				185					190				Lys	195
Leu	Ala	Met	Glu	185 Asp 200	Lys	His	Ile	Ile	190 Gln 205	Leu	Gln	Ser		195 Lys 210
Leu Glu	Ala Glu	Met Lys	Glu Asp	185 Asp 200 Gln 215	Lys · Leu	His Gln	Ile Val	Ile Leu	190 Gln 205 Val 220	Leu Ser	Gln Lys	Ser	Ile	195 Lys 210 Ser 225
Leu Glu Ile	Ala Glu Ile	Met Lys Glu	Glu Asp Glu	185 Asp 200 Gln 215 Leu 230	Lys Leu Glu	His Gln Lys	Ile Val Lys	Ile Leu Ile	190 Gln 205 Val 220 Val 235	Leu Ser Thr	Gln Lys Ala	Ser Gln Thr	Ile	195 Lys 210 Ser 225 Asn 240
Leu Glu Ile Asn	Ala Glu Ile Ser	Met Lys Glu Val	Glu Asp Glu Leu	Asp 200 Gln 215 Leu 230 Gln 245	Lys Leu Glu Lys	His Gln Lys Gln	Ile Val Lys Gln	Ile Leu Ile His	190 Gln 205 Val 220 Val 235 Asp 250	Leu Ser Thr	Gln Lys Ala Met	Ser Gln Thr	Ile Asn Val	195 Lys 210 Ser 225 Asn 240 Val 255
Leu Glu Ile Asn	Ala Glu Ile Ser Asn	Met Lys Glu Val Leu	Glu Asp Glu Leu Leu	Asp 200 Gln 215 Leu 230 Gln 245 Thr 260	Lys Leu Glu Lys Met	His Gln Lys Gln Met	Ile Val Lys Gln Ser	Ile Leu Ile His	190 Gln 205 Val 220 Val 235 Asp 250 Ser 265	Leu Ser Thr Leu Asn	Gln Lys Ala Met Ser	Ser Gln Thr Glu	Ile Asn Val	195 Lys 210 Ser 225 Asn 240 Val 255 Asp 270
Leu Glu Ile Asn Asn	Ala Glu Ile Ser Asn	Met Lys Glu Val Leu	Glu Asp Glu Leu Leu	Asp 200 Gln 215 Leu 230 Gln 245 Thr 260 Lys 275	Lys Leu Glu Lys Met	His Gln Lys Gln Met	Ile Val Lys Gln Ser	Ile Leu Ile His Thr	190 Gln 205 Val 220 Val 235 Asp 250 Ser 265 Ser 280	Leu Ser Thr Leu Asn	Gln Lys Ala Met Ser	Ser Gln Thr Glu Ala Asp	Ile Asn Val Thr	195 Lys 210 Ser 225 Asn 240 Val 255 Asp 270 Ala 285
Leu Glu Ile Asn Asn Pro	Ala Glu Ile Ser Asn Thr	Met Lys Glu Val Leu Val	Glu Asp Glu Leu Leu Ala	Asp 200 Gln 215 Leu 230 Gln 245 Thr 260 Lys 275 Ser 290	Lys Leu Glu Lys Met Glu Gly	His Gln Lys Gln Met Glu His	Ile Val Lys Gln Ser Gln Thr	Ile Leu Ile His Thr	190 Gln 205 Val 220 Val 235 Asp 250 Ser 265 Ser 280 Asn 295	Leu Ser Thr Leu Asn Phe	Gln Lys Ala Met Ser Arg	Ser Gln Thr Glu Ala Asp	Ile Asn Val Thr Lys	195 Lys 210 Ser 225 Asn 240 Val 255 Asp 270 Ala 285 Leu 300

Gly Ser Val Asp Phe Gln Arg Thr Trp Lys Glu Tyr Lys Val Gly 335 Phe Gly Asn Pro Ser Gly Glu Tyr Trp Leu Gly Asn Glu Phe Val 350 355 Ser Gln Leu Thr Asn Gln Gln Arg Tyr Val Leu Lys Ile His Leu Lys Asp Trp Glu Gly Asn Glu Ala Tyr Ser Leu Tyr Glu His Phe 380 385 390 Tyr Leu Ser Ser Glu Glu Leu Asn Tyr Arg Ile His Leu Lys Gly 400 405 Leu Thr Gly Thr Ala Gly Lys Ile Ser Ser Ile Ser Gln Pro Gly 415 420 Asn Asp Phe Ser Thr Lys Asp Gly Asp Asn Asp Lys Cys Ile Cys Lys Cys Ser Gln Met Leu Thr Gly Gly Trp Trp Phe Asp Ala Cys Gly Pro Ser Asn Leu Asn Gly Met Tyr Tyr Pro Gln Arg Gln Asn 465 Thr Asn Lys Phe Asn Gly Ile Lys Trp Tyr Tyr Trp Lys Gly Ser 475 Gly Tyr Ser Leu Lys Ala Thr Thr Met Met Ile Arg Pro Ala Asp 485 490 495

Phe

<210> 105 <211> 2076 <212> DNA

<213> Homo sapien

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<210> 106

<211> 419

<212> PRT

<213> Homo sapien

<400> 106

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Ala Ala Leu Leu Pro Gly Pro Arg Glu Ala Pro Ala Ala Ala Ala 20 25 30

Ala Phe Glu Ser Gly Leu Asp Leu Ser Asp Ala Glu Pro Asp Ala 35 40 40

Gly Glu Ala Thr Ala Tyr Ala Ser Lys Asp Leu Glu Glu Gln Leu 50 55 60

Arg Ser Val Ser Ser Val Asp Glu Leu Met Thr Val Leu Tyr Pro 65 70 75

Gln His Asn Arg Glu Gln Ala Asn Leu Asn Ser Arg Thr Glu Glu 95 100 105

Thr Ile Lys Phe Ala Ala Ala His Tyr Asn Thr Glu Ile Leu Lys 110 115 120

Ser Ile Asp Asn Glu Trp Arg Lys Thr Gln Cys Met Pro Arg Glu 125 130 135

Val Cys Ile Asp Val Gly Lys Glu Phe Gly Val Ala Thr Asn Thr  $140 \,$   $145 \,$   $150 \,$ 

Phe Phe Lys Pro Pro Cys Val Ser Val Tyr Arg Cys Gly Gly Cys 155 160 165

Cys Asn Ser Glu Gly Leu Gln Cys Met Asn Thr Ser Thr Ser Tyr 170 175 180

Leu Ser Lys Thr Leu Phe Glu Ile Thr Val Pro Leu Ser Gln Gly
185 190 190

Pro Lys Pro Val Thr Ile Ser Phe Ala Asn His Thr Ser Cys Arg
200 . 205 210

Cys Met Ser Lys Leu Asp Val Tyr Arg Gln Val His Ser Ile Ile 215 220 225

Arg Arg Ser Leu Pro Ala Thr Leu Pro Gln Cys Gln Ala Ala Asn 230 235 240

Lys Thr Cys Pro Thr Asn Tyr Met Trp Asn Asn His Ile Cys Arg 245 250 255

Cys Leu Ala Gln Glu Asp Phe Met Phe Ser Ser Asp Ala Gly Asp 260 265 270 Asp Ser Thr Asp Gly Phe His Asp Ile Cys Gly Pro Asn Lys Glu 275 280 285 Leu Asp Glu Glu Thr Cys Gln Cys Val Cys Arg Ala Gly Leu Arg 290 295 300 Pro Ala Ser Cys Gly Pro His Lys Glu Leu Asp Arg Asn Ser Cys 305 310 315 Gln Cys Val Cys Lys Asn Lys Leu Phe Pro Ser Gln Cys Gly Ala 325 330 Asn Arg Glu Phe Asp Glu Asn Thr Cys Gln Cys Val Cys Lys Arg 335 340 345 Thr Cys Pro Arg Asn Gln Pro Leu Asn Pro Gly Lys Cys Ala Cys 360 Glu Cys Thr Glu Ser Pro Gln Lys Cys Leu Leu Lys Gly Lys Lys 375 Phe His His Gln Thr Cys Ser Cys Tyr Arg Arg Pro Cys Thr Asn 385 Arg Gln Lys Ala Cys Glu Pro Gly Phe Ser Tyr Ser Glu Glu Val 395 400 Cys Arg Cys Val Pro Ser Tyr Trp Lys Arg Pro Gln Met Ser

<210> 107 <211> 2653

<212> DNA

<400> 107

<213> Homo sapien

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<212> PRT

<213> Homo sapien

<400> 108

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Thr Gln Met Ile Tyr Gln Gln His Gln Ser Trp Leu Arg Pro Val $50 \\ 55 \\ 60$ 

Leu Arg Ser Asn Arg Val Glu Tyr Cys Trp Cys Asn Ser Gly Arg
65 70 75

Ala Gln Cys His Ser Val Pro Val Lys Ser Cys Ser Glu Pro Arg 80 85 90

Cys Phe Asn Gly Gly Thr Cys Gln Gln Ala Leu Tyr Phe Ser Asp  $95\,$  100 105

Phe Val Cys Gln Cys Pro Glu Gly Phe Ala Gly Lys Cys Cys Glu 110 115 120

Ile Asp Thr Arg Ala Thr Cys Tyr Glu Asp Gln Gly Ile Ser Tyr 125 130 135

Arg Gly Thr Trp Ser Thr Ala Glu Ser Gly Ala Glu Cys Thr Asn 140 145 150

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Pro	Asp	Ala	Ile	Arg 170	Leu	Gly	Leu	Gly	Asn 175	His	Asn	Tyr	Cys	Arg 180
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Thr	His	Ser	Leu	Thr 230	Glu	Ser	Gly	Ala	Ser 235	Cys	Leu	Pro	Trp	Asn 240
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Leu	Arg	Gln	Tyr	Ser 305	Gln	Pro	Gln	Phe	Arg 310	Ile	Lys	Gly	Gly	Leu 315
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Lys	His	Arg	Arg	Ser 335	Pro	Gly	Glu	Arg	Phe 340	Leu	Cys	Gly	Gly	Ile 345
Leu	Ile	Ser	Ser	Cys 350	Trp	Ile	Leu	Ser	Ala 355	Ala	His	Cys	Phe	Gln 360
Glu	Arg	Phe	Pro	Pro 365	His	His	Leu	Thr	Val 370	Ile	Leu	Gly	Arg	Thr 375
Tyr	Arg	Val	Val	Pro 380	Gly	Glu	Glu	Glu	Gln 385	Lys	Phe	Glu	Val	Glu 390
Lys	Tyr	Ile	Val	His 395	Lys	Glu	Phe	Asp	Asp 400	Asp	Thr	Tyr	Asp	Asn 405
Asp	Ile	Ala	Leu	Leu 410	Gln	Leu	Lys	Ser	Asp 415	Ser	Ser	Arg	Cys	Ala 420
Gln	Glu	Ser	Ser	Val 425	Val	Arg	Thr	Val	Cys 430	Leu	Pro	Pro	Ala	Asp 435
Leu	Gln	Leu	Pro	Asp 440	Trp	Thr	Glu	Cys	Glu 445	Leu	Ser	Gly	Tyr	Gly 450
Lys	His	Glu	Ala	Leu	Ser	Pro	Phe	Tyr	Ser	Glu	Arg	Leu	Lys	Glu

455 460 465 Ala His Val Arg Leu Tyr Pro Ser Ser Arg Cys Thr Ser Gln His 470 475 Leu Leu Asn Arg Thr Val Thr Asp Asn Met Leu Cys Ala Gly Asp 485 . 490 495 Thr Arg Ser Gly Gly Pro Gln Ala Asn Leu His Asp Ala Cys Gln 500 505 Gly Asp Ser Gly Gly Pro Leu Val Cys Leu Asn Asp Gly Arg Met 515 525 Thr Leu Val Gly Ile Ile Ser Trp Gly Leu Gly Cys Gly Gln Lys

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Arg Gly Phe Gln Trp Val Thr Gly Asp Asn Asn Thr Ser Tyr Ser Arg Trp Ala Arg Leu Asp Leu Asn Gly Ala Pro Leu Cys Gly Pro Leu Cys Val Ala Val Ser Ala Ala Glu Ala Thr Val Pro Ser Glu 145 Pro Ile Trp Glu Glu Gln Cys Glu Val Lys Ala Asp Gly Phe 155 160 165 Leu Cys Glu Phe His Phe Pro Ala Thr Cys Arg Pro Leu Ala Val 170 175 Glu Pro Gly Ala Ala Ala Ala Val Ser Ile Thr Tyr Gly Thr 185 190 195 Pro Phe Ala Ala Arg Gly Ala Asp Phe Gln Ala Leu Pro Val Gly 205 Ser Ser Ala Ala Val Ala Pro Leu Gly Leu Gln Leu Met Cys Thr 215 225 Ala Pro Pro Gly Ala Val Gln Gly His Trp Ala Arg Glu Ala Pro 235

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Tyr	Gln	Cys	Gln	Pro 380	Leu	Asn	Gln	Thr	Ser 385	Tyr	Leu	Cys	Val	Cys 390
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Thr	Gln	Ala	Ser	Cys 425	Glu	Cys	Pro	Glu	Gly 430	Tyr	Ile	Leu	Asp	Asp 435
Gly	Phe	Ile	Cys	Thr 440	Asp	Ile	Asp	Glu	Cys 445	Glų	Asn	Gly	Gly	Phe 450
Cys	Ser	Gly	Val	Cys 455	His	Asn	Leu	Pro	Gly 460	Thr	Phe	Glu	Cys	Ile 465
Cys	Gly	Pro	Asp	Ser 470	Ala	Leu	Ala	Arg	His 475	Ile	Gly	Thr	Asp	Cys 480
Asp	Ser	Gly	Lys	Val 485	Asp	Gly	Gly	Asp	Ser 490	Gly	Ser	Gly	Glu	Pro 495
Pro	Pro	Ser	Pro	Thr 500	Pro	Gly	Ser	Thr	Leu 505	Thr	Pro	Pro	Ala	Val 510
				500					505				Ala Ala	510

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<213> Homo sapien

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<210> 112

<211> 295

<212> PRT

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Trp Lys Ser Thr Asn Phe Lys Thr Ile Leu Glu Trp Glu Pro Lys 50 55 60

Pro Val Asn Gln Val Tyr Thr Val Gln Ile Ser Thr Lys Ser Gly

65 70 75 Asp Trp Lys Ser Lys Cys Phe Tyr Thr Thr Asp Thr Glu Cys Asp Leu Thr Asp Glu Ile Val Lys Asp Val Lys Gln Thr Tyr Leu Ala 100 Arg Val Phe Ser Tyr Pro Ala Gly Asn Val Glu Ser Thr Gly Ser 110 Ala Gly Glu Pro Leu Tyr Glu Asn Ser Pro Glu Phe Thr Pro Tyr 130 135 Leu Glu Thr Asn Leu Gly Gln Pro Thr Ile Gln Ser Phe Glu Gln Val Gly Thr Lys Val Asn Val Thr Val Glu Asp Glu Arg Thr Leu Val Arg Arg Asn Asn Thr Phe Leu Ser Leu Arg Asp Val Phe Gly Lys Asp Leu Ile Tyr Thr Leu Tyr Tyr Trp Lys Ser Ser Ser Ser 190 Gly Lys Lys Thr Ala Lys Thr Asn Thr Asn Glu Phe Leu Ile Asp 205 210 Val Asp Lys Gly Glu Asn Tyr Cys Phe Ser Val Gln Ala Val Ile 220 Pro Ser Arg Thr Val Asn Arg Lys Ser Thr Asp Ser Pro Val Glu 230 235 240 Cys Met Gly Gln Glu Lys Gly Glu Phe Arg Glu Ile Phe Tyr Ile 250 Ile Gly Ala Val Val Phe Val Val Ile Ile Leu Val Ile Ile Leu 260 265 270

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<211> 2696

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275

Trp Lys Glu Asn Ser Pro Leu Asn Val Ser 290

280

295

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<210> 114

<211> 362

<212> PRT

<213> Homo sapien

# <400> 114

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Asp His Leu Phe Pro Pro Ser Leu Tyr Ile Phe Val Ile Gly Val
20 25 30

Gly Leu Pro Thr Asn Cys Leu Ala Leu Trp Ala Ala Tyr Arg Gln 35 40 45

Val Gln Gln Arg Asn Glu Leu Gly Val Tyr Leu Met Asn Leu Ser
50 55 60

Ile Ala Asp Leu Leu Tyr Ile Cys Thr Leu Pro Leu Trp Val Asp 65 70 75

Tyr Phe Leu His His Asp Asn Trp Ile His Gly Pro Gly Ser Cys

80 85 90 Lys Leu Phe Gly Phe Ile Phe Tyr Thr Asn Ile Tyr Ile Ser Ile 100 Ala Phe Leu Cys Cys Ile Ser Val Asp Arg Tyr Leu Ala Val Ala 115 His Pro Leu Arg Phe Ala Arg Leu Arg Arg Val Lys Thr Ala Val 130 Ala Val Ser Ser Val Val Trp Ala Thr Glu Leu Gly Ala Asn Ser Ala Pro Leu Phe His Asp Glu Leu Phe Arg Asp Arg Tyr Asn His Thr Phe Cys Phe Glu Lys Phe Pro Met Glu Gly Trp Val Ala Trp Met Asn Leu Tyr Arg Val Phe Val Gly Phe Leu Phe Pro Trp Ala 190 Leu Met Leu Ser Tyr Arg Gly Ile Leu Arg Ala Val Arg Gly Ser Val Ser Thr Glu Arg Gln Glu Lys Ala Lys Ile Lys Arg Leu 215 220 225 Ala Leu Ser Leu Ile Ala Ile Val Leu Val Cys Phe Ala Pro Tyr 235 His Val Leu Leu Ser Arg Ser Ala Ile Tyr Leu Gly Arg Pro 245 250 255 Trp Asp Cys Gly Phe Glu Glu Arg Val Phe Ser Ala Tyr His Ser 265 Ser Leu Ala Phe Thr Ser Leu Asn Cys Val Ala Asp Pro Ile Leu 275 280 285 Tyr Cys Leu Val Asn Glu Gly Ala Arg Ser Asp Val Ala Lys Ala 295 Leu His Asn Leu Leu Arg Phe Leu Ala Ser Asp Lys Pro Gln Glu 305 310 315 Met Ala Asn Ala Ser Leu Thr Leu Glu Thr Pro Leu Thr Ser Lys 325 Arg Asn Ser Thr Ala Lys Ala Met Thr Gly Ser Trp Ala Ala Thr Pro Pro Ser Gln Gly Asp Gln Val Gln Leu Lys Met Leu Pro Pro 355 Ala Gln

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<210> 116

<211> 426

<212> PRT

<213> Homo sapien

### <400> 116

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- Tyr Pro Gly Gly Ala Arg Asn Pro Met Ala Cys Asn Gly Ser Ala 20 25 30
- Ala Arg Gly His Phe Asp Pro Glu Asp Leu Asn Leu Thr Asp Glu
  35 40 45
- Ala Leu Arg Leu Lys Tyr Leu Gly Pro Gln Gln Thr Glu Leu Phe 50 55 60
- Met Pro Ile Cys Ala Thr Tyr Leu Leu Ile Phe Val Val Gly Ala 65 70 75
- Val Gly Asn Gly Leu Thr Cys Leu Val Ile Leu Arg His Lys Ala 80 85 90
- Met Arg Thr Pro Thr Asn Tyr Tyr Leu Phe Ser Leu Ala Val Ser 95 100 105
- Asp Leu Leu Val Leu Val Gly Leu Pro Leu Glu Leu Tyr Glu
  110 115 120
- Met Trp His Asn Tyr Pro Phe Leu Leu Gly Val Gly Gly Cys Tyr
  125
  130
  135
- Phe Arg Thr Leu Leu Phe Glu Met Val Cys Leu Ala Ser Val Leu 140 145 150
- Asn Val Thr Ala Leu Ser Val Glu Arg Tyr Val Ala Val Val His 155 160 165
- Pro Leu Gln Ala Arg Ser Met Val Thr Arg Ala His Val Arg Arg
  170 175 180
- Val Leu Gly Ala Val Trp Gly Leu Ala Met Leu Cys Ser Leu Pro

				185					190					195
Asn	Thr	Ser	Leu	His 200	Gly	Ile	Gln	Gln	Leu 205	His	Val	Pro	Cys	Arg 210
Gly	Pro	Val	Pro	Asp 215	Ser	Ala	Val	Cys	Met 220	Leu	Val	Arg	Pro	Arg 225
Ala	Leu	Tyr	Asn	Met 230	Val	Val	Gln	Thr	Thr 235	Ala	Leu	Leu	Phe	Phe 240
Cys	Leu	Pro	Met	Ala 245	Ile	Met	Ser	Val	Leu 250	Tyr	Leu	Leu	Ile	Gly 255
Leu	Arg	Leu	Arg	Arg 260	Glu	Arg	Leu	Leu	Leu 265	Met	Gln	Glu	Ala	Lys 270
Gly	Arg	Gly	Ser	Ala 275	Ala	Ala	Arg	Ser	Arg 280	Tyr	Thr	Cys	Arg	Leu 285
Gln	Gln	His	Asp	Arg 290	Gly	Arg	Arg	Gln	Val 295	Thr	Lys	Met	Leu	Phe 300
Val	Leu	Val	Val	Val 305	Phe	Gly	Ile	Cys	Trp 310	Ala	Pro	Phe	His	Ala 315
Asp	Arg	Val	Met	Trp 320	Ser	Val	Val	Ser	Gln 325	Trp	Thr	Asp	Gly	Leu 330
His	Leu	Ala	Phe	Gln 335	His	Val	His	Val	Ile 340	Ser	Gly	Ile	Phe	Phe 345
Tyr	Leu	Gly	Ser	Ala 350	Ala	Asn	Pro	Val	Leu 355	Tyr	Ser	Leu	Met	Ser 360
Ser	Arg	Phe	Arg	Glu 365	Thr	Phe	Gln	Glu	Ala 370	Leu	Cys	Leu	Gly	Ala 375
Cys	Cys	His	Arg	Leu 380	Arg	Pro	Arg	His	Ser 385	Ser	His	Ser	Leu	Ser 390
Arg	Met	Thr	Thr	Gly 395	Ser	Thr	Leu	Cys	Asp 400	Val	Gly	Ser	Leu	Gly 405
Ser	Trp	Val	His	Pro 410	Leu	Ala	Gly	Asn	Asp 415	Gly	Pro	Glu	Ala	Gln 420
Gln	Glu	Thr	Asp	Pro 425	Ser									

<210> 117 <211> 2512

<212> DNA

<213> Homo sapien

<400> 117

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gccctcctgc ctgcaggatc atgcccacca ccgtggacga tgtcctggag 200 catggaggg agtttcactt tttccagaag caaatgtttt tcctcttggc 250 tetgeteteg getacetteg egeceateta egtgggeate gtetteetgg 300 gcttcacccc tgaccaccgc tgccggagcc ccggagtggc cgagctgagt 350 ctgcgctgcg gctggagtcc tgcagaggaa ctgaactaca cggtgccggg 400 cccaggacct gcgggcgaag cctccccaag acagtgtagg cgctacgagg 450 tggactggaa ccagagcacc ttcgactgcg tggaccccct ggccagcctg 500 gacaccaaca ggagccgcct gccactgggc ccctgccggg acggctgggt 550 gtacgagacg cctggctcgt ccatcgtcac cgagtttaac ctggtatgtg 600 ccaactcctg gatgttggac ctattccagt catcagtgaa tgtaggattc 650 tttattggct ctatgagtat cggctacata gcagacaggt ttggccgtaa 700 gctctgcctc ctaactacag tcctcataaa tgctgcagct ggagttctca 750 tggccatttc cccaacctat acgtggatgt taatttttcg cttaatccaa 800 ggactggtca gcaaagcagg ctggttaata ggctacatcc tgattacaga 850 atttgttggg cggagatatc ggagaacagt ggggattttt taccaagttg 900 cctatacagt tgggctcctg gtgctagctg gggtggctta cgcacttcct 950 cactggaggt ggttgcagtt cacagttgct ctgcccaact tcttcttctt 1000 gctctattac tggtgcatac ctgagtctcc caggtggctg atctcccaga 1050 ataagaatgc tgaagccatg agaatcatta agcacatcgc aaagaaaaat 1100 ggaaaatctc tacccgcctc ccttcagcgc ctgagacttg aagaggaaac 1150 tggcaagaaa ttgaaccctt catttcttga cttggtcaga actcctcaga 1200 taaggaaaca tactatgata ttgatgtaca actggttcac gagctctgtg 1250 ctctaccagg gcctcatcat gcacatgggc cttgcaggtg acaatatcta 1300 cctggatttc ttctactctg ccctggttga attcccagct gccttcatga 1350 teatecteae categacege ateggaegee gttaceettg ggetgeatea 1400 aatatggttg caggggcagc ctgtctggcc tcagttttta tacctggtga 1450 tctacaatgg ctaaaaatta ttatctcatg cttgggaaga atggggatca 1500 caatggccta tgagatagtc tgcctggtca atgctgagct gtaccccaca 1550 ttcattagga atcttggcgt ccacatctgt tcctcaatgt gtgacattgg 1600 tggcatcatc acgccattcc tggtctaccg gctcactaac atctggcttg 1650

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<210> 118

<211> 555

<212> PRT

<213> Homo sapien

# <400> 118

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His Phe Phe Gln Lys Gln Met Phe Phe Leu Leu Ala Leu Leu Ser 20 25 30

Ala Thr Phe Ala Pro Ile Tyr Val Gly Ile Val Phe Leu Gly Phe
35 40 45

Thr Pro Asp His Arg Cys Arg Ser Pro Gly Val Ala Glu Leu Ser 50 55 60

Leu Arg Cys Gly Trp Ser Pro Ala Glu Glu Leu Asn Tyr Thr Val
65 70 75

Pro Gly Pro Gly Pro Ala Gly Glu Ala Ser Pro Arg Gln Cys Arg 80 85 90

Arg Tyr Glu Val Asp Trp Asn Gln Ser Thr Phe Asp Cys Val Asp

				95					100					105
Pro	Leu	Ala	Ser	Leu 110	Asp	Thr	Asn	Arg	Ser 115	Arg	Leu	Pro	Leu	Gly 120
Pro	Cys	Arg	Asp	Gly 125	Trp	Val	Tyr	Glu	Thr 130	Pro	Gly	Ser	Ser	Ile 135
Val	Thr	Glu	Phe	Asn 140	Leu	Val	Cys	Ala	Asn 145	Ser	Trp	Met	Leu	Asp 150
Leu	Phe	Gln	Ser	Ser 155	Val	Asn	Val	Gly	Phe 160	Phe	Ile	Gly	Ser	Met 165
Ser	Ile	Gly	Tyr	Ile 170	Ala	Asp	Arg	Phe	Gly 175	Arg	Lys	Leu	Cys	Leu 180
Leu	Thr	Thr	Val	Leu 185	Ile	Asn	Ala	Ala	Ala 190	Gly	Val	Leu	Met	Ala 195
Ile	Ser	Pro	Thr	Tyr 200	Thr	Trp	Met	Leu	Ile 205	Phe	Arg	Leu	Ile	Gln 210
Gly	Leu	Val	Ser	Lys 215	Ala	Gly	Trp	Leu	Ile 220	Gly	Tyr	Ile	Leu	Ile 225
Thr	Glu	Phe	Val	Gly 230	Arg	Arg	Tyr	Arg	Arg 235	Thr	Val	Gly	Ile	Phe 240
Tyr	Gln	Val	Ala	Tyr 245	Thr	Val	Gly	Leu	Leu 250	Val	Leu	Ala	Gly	Val 255
Ala	Tyr	Ala	Leu	Pro 260	His	Trp	Arg	Trp	Leu 265	Gln	Phe	Thr	Val	Ala 270
Leu	Pro	Asn	Phe	Phe 275	Phe	Leu	Leu	Tyr	Tyr 280	Trp	Cys	Ile	Pro	Glu 285
Ser	Pro	Arg	Trp	Leu 290	Ile	Ser	Gln	Asn	Lys 295	Asn	Ala	Glu	Ala	Met 300
Arg	Ile	Ile	Lys	His 305	Ile	Ala	Lys	Lys	Asn 310	Gly	Lys	Ser	Leu	Pro 315
Ala	Ser	Leu	Gln	Arg 320	Leu	Arg	Leu	Glu	Glu 325	Glu	Thr	Gly	Lys	Lys 330
Leu	Asn	Pro	Ser	Phe 335	Leu	Asp	Leu	Val	Arg 340	Thr	Pro	Gln	Ile	Arg 345
Lys	His	Thr	Met	Ile 350	Leu	Met	Tyr	Asn	Trp 355	Phe	Thr	Ser	Ser	Val 360
Leu	Tyr	Gln	Gly	Leu 365	Ile	Met	His	Met	Gly 370	Leu	Ala	Gly	Asp	Asn 375
Ile	Tyr	Leu	Asp	Phe 380	Phe	Tyr	Ser	Ala	Leu 385	Val	Glu	Phe	Pro	Ala 390
Ala	Phe	Met	Ile	Ile 395	Leu	Thr	Ile	Asp	Arg 400	Ile	Gly	Arg	Arg	Tyr 405

Pro Trp Ala Ala Ser Asn Met Val Ala Gly Ala Ala Cys Leu Ala 410 415 420 Ser Val Phe Ile Pro Gly Asp Leu Gln Trp Leu Lys Ile Ile Ile 430 Ser Cys Leu Gly Arg Met Gly Ile Thr Met Ala Tyr Glu Ile Val 445 450 Cys Leu Val Asn Ala Glu Leu Tyr Pro Thr Phe Ile Arg Asn Leu 455 460 Gly Val His Ile Cys Ser Ser Met Cys Asp Ile Gly Gly Ile Ile Thr Pro Phe Leu Val Tyr Arg Leu Thr Asn Ile Trp Leu Glu Leu 490 495 Pro Leu Met Val Phe Gly Val Leu Gly Leu Val Ala Gly Gly Leu 505 510 Val Leu Leu Pro Glu Thr Lys Gly Lys Ala Leu Pro Glu Thr 515 520 525 Ile Glu Glu Ala Glu Asn Met Gln Arg Pro Arg Lys Asn Lys Glu 530 535 Lys Met Ile Tyr Leu Gln Val Gln Lys Leu Asp Ile Pro Leu Asn 545 550 555

<210> 119

<211> 5431

<212> DNA

<213> Homo sapien

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<211> 1533

<212> PRT

<213> Homo sapien

<400> 120

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Ile Ser Val His Gly Gly Leu Gln Asn Phe Glu Met Gln Pro Lys  $^{\circ}$  35 40 45

Leu Lys Gln Val Phe Gly Lys Gly Leu Ile Lys Ala Ala Met Thr 50 55 60

Thr Gly Ala Trp Ile Phe Thr Gly Gly Val Ser Thr Gly Val Ile 65 70 75

Ser His Val Gly Asp Ala Leu Lys Asp His Ser Ser Lys Ser Arg 80 85 90

Gly Arg Val Cys Ala Ile Gly Ile Ala Pro Trp Gly Ile Val Glu 95 100 105

Asn Lys Glu Asp Leu Val Gly Lys Asp Val Thr Arg Val Tyr Gln \$110\$ \$120\$

Thr Met Ser Asn Pro Leu Ser Lys Leu Ser Val Leu Asn Asn Ser 125 130 135

His Thr His Phe Ile Leu Ala Asp Asn Gly Thr Leu Gly Lys Tyr
140 145 150

Gly Ala Glu Val Lys Leu Arg Arg Leu Leu Glu Lys His Ile Ser 155 160 165

Leu Gln Lys Ile Asn Thr Arg Leu Gly Gln Gly Val Pro Leu Val
170 175 180

Gly Leu Val Val Glu Gly Gly Pro Asn Val Val Ser Ile Val Leu 185 190 195

Glu Tyr Leu Gln Glu Glu Pro Pro Ile Pro Val Val Ile Cys Asp 200 205 210

Gly Ser Gly Arg Ala Ser Asp Ile Leu Ser Phe Ala His Lys Tyr 215 220 225

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Leu	Val	Thr	Ile	Gln 245	Lys	Thr	Phe	Asn	Tyr 250	Asn	Lys	Ala	Gln	Ser 255
His	Gln	Leu	Phe	Ala 260	Ile	Ile	Met	Glu	Cys 265	Met	Lys	Lys	Lys	Glu 270
Leu	Val	Thr	Val	Phe 275	Arg	Met	Gly	Ser	Glu 280	Gly	Gln	Gln	Asp	Ile 285
Glu	Met	Ala	Ile	Leu 290	Thr	Ala	Leu	Leu	Lys 295	Gly	Thr	Asn	Val	Ser 300
Ala	Pro	Asp	Gln	Leu 305	Ser	Leu	Ala	Leu	Ala 310	Trp	Asn	Arg	Val	Asp 315
Ile	Ala	Arg	Ser	Gln 320	Ile	Phe	Val	Phe	Gly 325	Pro	His	Trp	Pro	Pro 330
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Pro	Pro	Asn	Thr	Leu 440	His	Leu	Leu	Val	Arg 445	Asp	Val	Lys	Lys	Ser 450
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Pro	Ala	Lys	Gly	Lys 515	Lys	Lys	Lys	Lys	Lys 520	Lys	Lys	Glu	Glu	Glu 525

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Lys	Met	Ala	Val	Phe 560	Leu	Trp	Gln	Arg	Gly 565	Glu	Glu	Ser	Met	Ala 570
Lys	Ala	Leu	Val	Ala 575	Cys	Lys	Leu	Tyr	Lys 580	Ala	Met	Ala	His	Glu 585
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Gln	Ser	Tyr	Lys	His 620	Asp	Glu	Gln	Ile	Ala 625	Met	Lys	Leu	Leu	Thr 630
Tyr	Glu	Leu	Lys	Asn 635	Trp	Ser	Asn	Ser	Thr 640	Cys	Leu	Lys	Leu	Ala 645
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Asn	Pro	Gly	Leu	Lys 680	Val	Ile	Met	Gly	Ile 685	Leu	Leu	Pro	Pro	Thr 690
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Cys	Glu	Phe	Tyr	Asn 755	Ala	Pro	Ile	Val	Lys 760	Phe	Trp	Phe	Tyr	Thr 765
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Tyr	Gly	Arg	Val	Ile 860	Tyr	Cys	Val	Asp	Ile 865	Ile	Phe	Trp	Tyr	Ile 870
Arg	Val	Leu	Asp	Ile 875	Phe	Gly	Val	Asn	Lys 880	Tyr	Leu	Gly	Pro	Tyr 885
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His	Phe	Arg		Lys 100	Glu	Asp	Glu	Gln 1	Gln 105	Ser	Ser	Ser	_	Glu 110
Arg	Ile	Arg		Thr 115	Ser	Glu	Arg	Val 1	Glu 120	Asn	Met	Ser		Arg 125
Leu	Glu	Glu		Asn .130	Glu	Arg	Glu	Thr 1	Phe 135	Met	Lys	Thr		Leu 140

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Val Ala Glu Ala Ile Pro Arg Ile Pro Arg Leu Ser Leu Thr Ile 1445 1450 1455

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Gln Thr Leu Gly Phe Pro Ser Leu Arg Ser Lys Ser Leu His Gly 1475 1480 1485

His Pro Arg Asn Val Lys Ser Ile Gln Gly Lys Leu Asp Arg Ser 1490 1495 1500

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<212> DNA

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<212> PRT

<213> Homo sapien

<400> 122

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Asn Ala Leu Thr Pro Pro Pro Thr Thr Pro Glu Trp Val Lys Phe 80 85 90

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Ala Ile Leu Cys Phe Leu Ala Tyr Gly Ile Gln Ala Ala Met Glu 110 115 120

Asp Glu Pro Ser Asn Asp Asn Leu Tyr Leu Gly Val Val Leu Ala 125 130 135

Ala Val Val Ile Val Thr Gly Cys Phe Ser Tyr Tyr Gln Giu Ala 140 145 150

Lys Ser Ser Lys Ile Met Asp Ser Phe Lys Asn Met Val Pro Gln 155 160 165

Gln Ala Leu Val Ile Arg Glu Gly Glu Lys Met Gln Ile Asn Ala 170 175 180

Glu Glu Val Val Gly Asp Leu Val Glu Val Lys Gly Gly Asp 185 190 195

Arg Val Pro Ala Asp Leu Arg Ile Ile Ser Ser His Gly Cys Lys
200 205 210

Val Asp Asn Ser Ser Leu Thr Gly Glu Ser Glu Pro Gln Thr Arg 215 220 225

Ser Pro Glu Phe Thr His Glu Asn Pro Leu Glu Thr Arg Asn Ile 230 235 240

Cys Phe Phe Ser Thr Asn Cys Val Glu Gly Thr Ala Arg Gly Ile \$245\$ \$250\$

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Ile	Glu	His	Phe	Ile 290	Gln	Leu	Ile	Thr	Gly 295	Val	Ala	Val	Phe	Leu 300
Gly	Val	Ser	Phe	Phe 305	Val	Leu	Ser	Leu	Ile 310	Leu	Gly	Tyr	Ser	Trp 315
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Pro	Glu	Gly	Leu	Leu 335	Ala	Thr	Val	Thr	Val 340	Cys	Leu	Thr	Leu	Thr 345
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